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(57) Abstract

Tumors attract blood vessels in order to grow by a process called angiogenesis. The relative quantity of stimulators and inhibitors is an important determining factor for the initiation of angiogenesis. Thrombospondins-1 and -2 are adhesive glycoproteine that have the ability to inhibit aniogenesis. This inhibiting activity has been mapped to the type 1 repeats of TSP-1 and TSP-2. The invention includes chimeric proteins that contain anti-angiogenic portions of TSP-1, TSP-2, endostatin, angiostatin, platelet factor 4, or prolactin, linked to a portion of the N-terminal region of human cartilage oligomeric matrix protein (COMP) that allows formation of pentamers. Also described herein are the nucleic acid molecules, vectors, and host cells for expressing and producing these chimeric proteins. Further embodiments of the invention include methods to treat humans or other mammals with anti-angiogenic proteins to reduce tumor size or rate of growth. Since the type 1 repeat region of TSP-1 and TSP-2 reportedly inhibits HIV infection, chimeric proteins comprising these repeats may also be used for this purpose, as well as to inhibit angiogenesis.

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COMP/TSP-1, COMP/TSP-2 AND OTHER CHIMERIC PROTEINS

RELATED APPLICATIONS

This application claims the benefit of U.S. Provisional Application No. 60/118,053 filed February 1, 1999, the entire teachings of which are incorporated herein by reference.

BACKGROUND OF THE INVENTION

Thrombospondins are a family of calcium-binding multifunctional glycoproteins that are secreted by various cell types and are developmentally regulated components of the extracellular matrix (Bornstein, P., FASEB J., 6:3290-3299, 1992; Bornstein, P., J. Cell Biol., 130:503-506, 1995). Among their functions are modulating cell attachment, migration and proliferation.

One member of this family, cartilage oligomeric matrix protein (COMP) is a pentamer in which multimerization appears to be directed by α-helical segments situated (in the amino acid sequence) either before or after the cysteine residues that form the interchain disulfide bonds. COMP has been purified (Prochownik, E.V. et al., J. Cell Biol. 109:843-852 (1989)). Individuals affected with pseudoachondroplasia, who have considerably shortened stature as a result of premature cessation of bone growth, have been shown to have mutations in exon 17B of the COMP protein (Nature Genetics 10:325-329 (1995)).

In vitro assays have shown that platelet thrombospondin-1 is involved in thrombosis, fibrinolysis, wound healing, inflammation, tumor cell metastasis and angiogenesis. The major form of thrombospondin secreted by platelets and endothelial cells is TSP-1. Thrombospondin-1 (TSP-1) is an angiogenesis inhibitor that decreases tumor growth. Thrombospondin- 2 (TSP-2) is a related glycoprotein of similar structure and properties.

The thrombospondin type 1 repeats (TSRs; also "repeat regions" herein) have been shown to inhibit angiogenesis and HIV infection. However, other portions of the proteins have been shown to have a positive effect on endothelial cell

WO 00/44908

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growth. Thromobospondin-1 and -2 are similar in terms of their molecular architecture. Thrombospondin-1 and thrombospondin-2 each have three copies of the TSR. TSP-1 and TSP-2 are trimeric molecules. Thus, each fully assembled protein contains nine TSRs.

Whereas TSP-1 and TSP-2 are antiangiogenic, these proteins contain other domains that have additional activities that diminish the antiangiogenic activity. The isolated TSRs are more potent inhibitors of angiogenesis than the native molecules.

The ingrowth of new capillary networks into developing tumors is essential for the progression of cancer. Thus, the development of pharmaceuticals that inhibit the process of angiogenesis is an important therapeutic goal. As pointed out in a review by Folkman (Folkman, J., *Proc. Natl. Acad. Sci. USA 95*: 9064-9066, 1998), antiangiogenic therapy has little toxicity, does not require the therapeutic agent to enter tumor cells or cross the blood-brain barrier, controls tumor growth independently of growth of tumor cell heterogeneity, and does not induce drug resistance.

SUMMARY OF THE INVENTION

The invention includes chimeric proteins comprising: (1) a chimeric protein comprising the second and third type 1 repeats of human TSP-1, and which may also comprise the procollagen homology region of TSP-1; (2) a chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the second and third type 1 repeats of human TSP-1; (3) a chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the second and third type 1 repeats of human TSP-1, but not the TGF- β activation region of human TSP-1; (4) a chimeric protein comprising the multimerization domain of human COMP, the procollagen region, and the first, second, and third type 1 repeats of human TSP-1; (5) a chimeric protein comprising the three type 1 repeats of human TSP-2, and which may also comprise the procollagen homology region of TSP-2; (6) a chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP,

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and the three type 1 repeats of human TSP-2; and (7) variants of any of the above having anti-angiogenic activity. The invention further includes isolated nucleic acids encoding any of the above chimeric proteins, vectors comprising these nucleic acids, and host cells comprising any of said vectors. The chimeric proteins can be produced in host cells and used in methods for the treatment of a disease or medical condition characterized by abnormal or undesirable proliferation of blood vessels, such as that occurring in tumor growth.

BRIEF DESCRIPTION OF THE DRAWINGS

Figure 1 is a representation of the amino acid sequence of human TSP-1 (SEQ ID NO: 1). The type 1 repeats of TSP-1 are, as illustrated here, 1) amino acids 361-416; 2) amino acids 417-473; and 3) amino acids 474-530.

Figure 2 is a representation of the amino acid sequence of human TSP-2 (SEQ ID NO: 2). The type 1 repeats of TSP-2 are, as illustrated here, 1) amino acids 381-436; 2) amino acids 437-493; and 3) amino acids 494-550.

Figure 3 is a representation of the amino acid sequence of human COMP (SEQ ID NO: 3). The type 2 repeats of COMP are, as illustrated here, 1) amino acids 89-128; 2) amino acids 129-181; 3) amino acids 182-226; and 4) amino acids 227-268

Figures 4A and 4B together are a representation of the DNA sequence (SEQ ID NO: 4) of gene encoding a human COMP/TSP-1 chimeric protein and the amino acid sequence (SEQ ID NO: 5) of a human COMP/TSP-1 chimeric protein encoded by the DNA sequence above it.

Figure 5A and 5B together are a representation of the DNA sequence (SEQ ID NO: 6) of a gene encoding a human COMP/TSP-2 chimeric protein and the amino acid sequence (SEQ ID NO: 7) of a human COMP/TSP-2 chimeric protein encoded by the DNA sequence above it.

Figure 6 is a schematic representation of a few of the chimeric protein embodiments of the invention.

Figure 7 is a graph showing tumor volume (mm³) at 7, 14 and 21 days in the experiment described in Example 3, in which mice were injected with an unaltered

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(control) vector, pNeo (filled diamonds) or with an expression vector encoding COMP/TSP-1 chimeric protein (filled squares).

DETAILED DESCRIPTION OF THE INVENTION

Described herein is a protein that has the functional activity of the TSR but not other activities associated with TSP-1 or TSP-2, and is assembled into a multimeric structure. One embodiment of the invention is a chimeric protein that comprises the TSRs from TSP-1 or TSP-2 and the multimer assembly region of human cartilage oligomeric matrix protein (COMP), using a portion of the aminoterminal end. Other portions of TSP-1 or TSP-2 can be incorporated into the chimeric protein, such as the procollagen homology region of TSP-1 and/or TSP-2. The last two TSRs of TSP-1 are preferably used because the first TSR has the ability to activate transforming growth factor β (TGF- β), which stimulates tumor growth. The COMP assembly domain spontaneously forms a 5-stranded α -helical domain, allowing for the use of the COMP domain as a tool for pentamerization.

Thus, the COMP/TSP-1 construct contains the region for multimerization, the first type 2 repeat of human COMP (construct encodes amino acids 1-128) and the second and third TSRs of human TSP-1 (construct encodes amino acids 417-530). See the Table for active sequences of TSP-1 (taken from chapter 2, "The Primary Structure of the Thrombospondins" In *The Thrombospondin Gene Family* (J.C. Adams *et al.*, eds.) Springer-Verlag, Heidelberg (1995)). The assembled protein is a pentamer containing 10 copies of the TSR. Thus, COMP/TSP-1 and COMP/TSP-2 are expected to be more active than TSP-1 and TSP-2. COMP/TSP-1 and COMP/TSP-2 are expected to be correctly folded and multimeric so that they better mimic the natural proteins than peptides that are based on the TSR sequence.

The first type 2 repeat of COMP includes amino acid residues 73-130, based on the genomic sequence. The amount of COMP sequence at the 3' end can be increased or decreased to maximize activity. For example, two or more type 2 repeats of COMP can be included if moving the type 1 repeats of TSP-1 or TSP-2 farther out on the arms of the expressed protein increases its activity. Alternatively, "spacer" sequence not naturally occurring in COMP or in TSP-1 or TSP-2 can be

PCT/US00/02482

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added. The COMP/TSP-2 construct contains the same region of COMP and the three TSRs of human TSP-2 (construct encodes amino acids 381-550). When it is assembled to a pentamer this chimeric protein will contain 15 TSRs. Because these proteins are derived from portions of human proteins, they should not be immunogenic in humans.

Table: Active Regions of Interest Within Thrombospondin-1

Domain	Sequence	Function
Procollagen	NGVQYRN (SEQ ID NO: 8)	Anti-angiogenesis
homology		
Type 1 repeats	CSVTCG (SEQ ID NO: 9)	Cell binding
	WSXWSXW (SEQ ID NO: 10)	Heparin binding
	GGWSHW (SEQ ID NO: 11)	TGF-β and Fibronectin
		binding
	RFK	TGF-β activation
	SPWDICSVTCGGGVQKRSR	Anti-angiogenesis
	(SEQ ID NO: 12)	
Type 2 repeats	DVDEC(X) ₆ C(X) ₈ CENTDPGYNCLPC	Calcium binding
	(SEQ ID NO: 13)	

In one aspect, the invention comprises polynucleotides or nucleic acid molecules that encode chimeric proteins having portions whose amino acid sequences are derived from human TSP-1. By the genomic structure, the type 1 repeats of TSP-1 are amino acid residues 359-414 (first), amino acid residues 415-473 (second), and 474-531 (third). In one case, the chimeric protein encoded by the polynucleotides of the invention comprises the second and third type 1 repeats of human TSP-1. Such a chimeric protein may also comprise the procollagen homology region and the first type 1 repeat of TSP-1. If amino acid sequences that 20 activate $TGF-\beta$ are included in the product protein, and are found to reduce antiangiogenic activity, the RFK sequence can be mutated (to QFK, for example) to a

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sequence that does not activate TGF- β , by appropriate manipulations of the nucleic acid molecule or construct encoding the chimeric proteins. In another case, the chimeric proteins encoded by the polynucleotides of the invention are variants of the immediately aforementioned chimeric protein which have activity that is similar in quality and quantity (for example, plus or minus one order of magnitude in an assay) to the anti-angiogenic activity of the protein whose amino acid sequence is represented in Figures 4A and 4B. In another case, the chimeric proteins encoded by polynucleotides of the invention comprise the second and third type 1 repeats of human TSP-1, the multimerization domain of human COMP, and the first type 2 repeat of human COMP. In another case, the chimeric proteins encoded by the polynucleotides of the invention are variants of the immediately aforementioned chimeric protein which have activity that is similar in quality and quantity to the anti-angiogenic activity of the protein whose amino acid sequence is represented in Figures 4A and 4B.

In one aspect, the invention comprises polynucleotides or nucleic acid molecules that encode chimeric proteins having portions whose amino acid sequences are derived from human TSP-2. The genomic structure of the human TSP-2 gene, which would provide one way to define the boundaries of the repeats, has not been determined. In one case, the chimeric protein encoded by the polynucleotides of the invention comprises the three type 1 repeats of human TSP-2. In another case, the chimeric proteins encoded by the polynucleotides of the invention are variants of the immediately aforementioned chimeric proteins which have activity that is similar in quality and quantity to the anti-angiogenic activity of the protein whose amino acid sequence is represented in Figures 5A and 5B. In another case, the chimeric protein encoded by polynucleotides of the invention comprises the three type 1 repeats of human TSP-2, and the multimerization domain of human COMP. In another case, the chimeric proteins encoded by the polynucleotides of the invention are variants of the immediately aforementioned chimeric protein which have activity that is similar in quality and quantity to the anti-angiogenic activity of the protein whose amino acid sequence is represented in Figures 5A and 5B.

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The polynucleotides of the invention can be made by recombinant methods, can be made synthetically, can be replicated by enzymes in *in vitro* (e.g., PCR) or *in vivo* systems (e.g., by suitable host cells, when inserted into a vector appropriate for replication within the host cells), or can be made by a combination of methods. The polynucleotides of the invention can include DNA and its RNA counterpart.

As used herein, "nucleic acid," "nucleic acid molecule," "oligonucleotide" and "polynucleotide" include DNA and RNA and chemical derivatives thereof, including phosphorothicate derivatives and RNA and DNA molecules having a radioactive isotope or a chemical adduct such as a fluorophore, chromophore or biotin (which can be referred to as a "label"). The RNA counterpart of a DNA is a polymer of ribonucleotide units, wherein the nucleotide sequence can be depicted as having the base U (uracil) at sites within a molecule where DNA has the base T (thymidine).

Isolated nucleic acid molecules or polynucleotides can be purified from a natural source or can be made recombinantly. Polynucleotides referred to herein as "isolated" are polynucleotides purified to a state beyond that in which they exist in cells. They include polynucleotides obtained by methods described herein, similar methods or other suitable methods, and also include essentially pure polynucleotides produced by chemical synthesis or by combinations of biological and chemical methods, and recombinant polynucleotides that have been isolated. The term "isolated" as used herein for nucleic acid molecules, indicates that the molecule in question exists in a physical milieu distinct from that in which it occurs in nature. For example, an isolated polynucleotide may be substantially isolated with respect to the complex cellular milieu in which it naturally occurs, and may even be purified essentially to homogeneity, for example as determined by agarose or polyacrylamide gel electorphoresis or by A_{260}/A_{280} measurements, but may also have further cofactors or molecular stabilizers (for instance, buffers or salts) added.

The invention further comprises the polypeptides encoded by the isolated nucleic acid molecules of the invention. Thus, for example, the invention relates to fusion proteins, comprising a portion of TSP-1 which comprises the second and third type 1 repeats, linked to a second moiety not occurring in TSP-1 as found in

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nature. In an analogous manner, the invention relates also to fusion proteins, comprising TSP-2 or a functional portion thereof such as one or more repeat regions as a first moiety, linked to second moiety not occurring in TSP-2 as found in nature. The second moiety can be an amino acid, peptide or polypeptide, and can have enzymatic or binding activity of its own. The first moiety can be in an N-terminal location, C-terminal location or internal to the fusion protein. In one embodiment, the fusion protein comprises the portion of human TSP-1 described immediately above, or human TSP-2 or a portion thereof as the first moiety, and a second moiety comprising a linker sequence and an affinity ligand.

Another aspect of the invention relates to a method of producing a chimeric protein of the invention, or a variant thereof, and to expression systems and host cells containing a vector appropriate for expression of a chimeric protein of the invention. Variants of the chimeric protein include those having amino acid sequences that differ from those sequences in Figures 4A and 4B, and Figures 5A and 5B, wherein those variants have several, such as 5 to 10, 1 to 5, or 3, 2 or 1 amino acids substituted, deleted, or added, in any combination, compared to the sequences in Figures 4A and 4B and Figures 5A and 5B. In one embodiment, variants have silent substitutions, additions and deletions that do not alter the properties and activities of the chimeric protein. Variants can also be modified polypeptides in which one or more amino acid residues are modified, and mutants comprising one or more modified residues.

Proteins and polypeptides described herein can be assessed for their angiogenic activity by using an assay such as those described in Tolsma, S.S. et al., J. Cell Biol. 122(2):497-511 (1993), one which measures the migration of bovine adrenal capillary endothelial cells in culture, and one which tests migration of cells into a sponge containing an agent to be tested for activity. A further test for angiogenesis, which can also be adapted also to test anti-angiogenesis activity, is described in Polyerini, P.J. et al., Methods. Enzymol. 198:440-450 (1991).

Cells that express such a chimeric protein or a variant thereof can be made and maintained in culture, under conditions suitable for expression, to produce protein for isolation. These cells can be procaryotic or eucaryotic. Examples of

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procaryotic cells that can be used for expression (as "host cells"; "cell" including herein cells of tissues, cell cultures, cell strains and cell lines) include *Escherichia coli, Bacillus subtilis* and other bacteria. Examples of eucaryotic cells that can be used for expression include yeasts such as *Saccharomyces cerevisiae*,

5 Schizosaccharomyces pombe, Pichia pastoris and other lower eucaryotic cells, and cells of higher eucaryotes such as those from insects and mammals. Suitable cells of mammalian origin include primary cells, and cell lines such as CHO, HeLa, 3T3, BHK, COS, 293, and Jurkat cells. Suitable cells of insect origin include primary cells, and cell lines such as SF9 and High five cells. (See, e.g., Ausubel, F.M. et al., eds. Current Protocols in Molecular Biology, Greene Publishing Associates and John Wiley & Sons Inc., (containing Supplements up through 1998)).

In one embodiment, host cells that produce a recombinant chimeric protein, variant, or portions thereof can be made as follows. A gene encoding a chimeric protein described herein can be inserted into a nucleic acid vector, e.g., a DNA vector, such as a plasmid, virus or other suitable replicon (including vectors suitable for use in gene therapy, such as those derived from adenovirus or others; see, for example Xu, M. et al., Molecular Genetics and Metabolism 63:103-109, 1998) can be present in a single copy or multiple copies, or the gene can be integrated in a host cell chromosome. A suitable replicon or integrated gene can contain all or part of the coding sequence for the protein or variant, operably linked to one or more expression control regions whereby the coding sequence is under the control of transcription signals and linked to appropriate translation signals to permit translation. The vector can be introduced into cells by a method appropriate to the type of host cells (e.g., transformation, electroporation, infection). For expression from the gene, the host cells can be maintained under appropriate conditions (e.g., in the presence of inducer, normal growth conditions, etc.). Proteins or polypeptides thus produced can be recovered (e.g., from the cells, the periplasmic space, culture medium) using suitable techniques.

The invention also relates to isolated proteins or polypeptides encoded by nucleic acids of the present invention. Isolated proteins can be purified from a natural source or can be made recombinantly. Proteins or polypeptides referred to

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herein as "isolated" are proteins or polypeptides purified to a state beyond that in which they exist in cells and include proteins or polypeptides obtained by methods described herein, similar methods or other suitable methods, and also include essentially pure proteins or polypeptides, proteins or polypeptides produced by chemical synthesis or by combinations of biological and chemical methods, and recombinant proteins or polypeptides which are isolated. Thus, the term "isolated" as used herein, indicates that the polypeptide in question exists in a physical milieu distinct from the cell in which its biosynthesis occurs. For example, an isolated COMP/TSP-1 or COMP/TSP-2 chimeric protein may be purified essentially to homogeneity, for example as determined by PAGE or column chromatography (for 10 example, HPLC), but may also have further cofactors or molecular stabilizers added to the purified protein to enhance activity. In one embodiment, proteins or polypeptides are isolated to a state at least about 75% pure; more preferably at least about 85% pure, and still more preferably at least about 95% pure, as determined by Coomassie blue staining of proteins on SDS-polyacrylamide gels.

Chimeric or fusion proteins can be produced by a variety of methods. For example, a chimeric protein can be produced by the insertion of a TSP gene or portion thereof into a suitable expression vector, such as Bluescript SK +/-(Stratagene), pGEX-4T-2 (Pharmacia), pET-15b, pET-20b(+) or pET-24(+) (Novagen). The resulting construct can be introduced into a suitable host cell for expression. Upon expression, chimeric protein can be purified from a cell lysate by means of a suitable affinity matrix (see e.g., Current Protocols in Molecular Biology (Ausubel, F.M. et al., eds., Vol. 2, pp. 16.4.1-16.7.8, containing supplements up through Supplement 44, 1998).

Polypeptides of the invention can be recovered and purified from cell cultures by well-known methods. The recombinant protein can be purified by ammonium sulfate precipitation, heparin-Sepharose affinity chromatography, gel filtration chromatography and/or sucrose gradient ultracentrifugation using standard techniques. Further methods that can be used for purification of the polypeptide include ethanol precipitation, acid extraction, anion or cation exchange chromatography, phosphocellulose chromatography, hydrophobic interaction

chromatography, affinity chromatography, hydroxylapatite chromatography and high performance liquid chromatography. Known methods for refolding protein can be used to regenerate active conformation if the polypeptide is denatured during isolation or purification.

hybrid proteins can be applied more broadly to produce polynucleotides, and vectors and host cells comprising such polynucleotides, wherein the polynucleotides encode COMP/endostatin, COMP/angiostatin, COMP/platelet factor 4, or COMP/prolactin, for example. In each case, a portion of a polynucleotide known to encode full-length human endostatin, angiostatin, platelet factor 4 (GenBank Accession No. M25897) or prolactin (GenBank Accession No. V00566), can be chosen for cloning into a COMP cDNA as illustrated herein for COMP/TSP-1 and COMP/TSP-2 DNA constructs. Thus, the invention also includes COMP/endostatin, COMP/angiostatin, COMP/platelet factor 4, and COMP/prolactin chimeric proteins encoded by such nucleic acid constructs. See Figure 6 for a schematic representation of the structure of COMP/endostatin.

In addition, a portion of the endostatin, angiostatin, platelet factor 4 or prolactin coding regions, wherein that portion encodes a polypeptide having antiangiogenic activity, can be added to or incorporated into a DNA construct encoding COMP/TSP-1, such that a TSP-1-derived polypeptide and a polypeptide derived from endostatin, angiostatin, platelet factor 4 or prolactin are produced fused together in tandem on the same "arm" of the "5-armed" COMP-multimerized pentamer. Different expression constructs can be introduced into the same host cells such that two or more chimeric protein "arms" of different types (e.g.,

25 COMP/angiostatin and COMP/TSP-1 or COMP/TSP-2) are joined at the COMP multimerization domain.

Chimeric protein antiangiogenic agents can be used, for example, after surgery or radiation to prevent recurrence of metastases, in combination with conventional chemotherapy, immunotherapy, or various types of gene therapy not necessarily directed against angiogenesis.

Construction of COMP/TSP-1P Expression Vectors

Expression vectors that can be used to produce COMP/TSP-1P, a chimeric protein that includes the procollagen homology region (see Figure 6), can be produced from two distinct cDNAs. The COMP portion is identical to that in the 5 Examples described herein. For TSP-1, a new forward primer (GAT GAC GTC ACT GAA GAG AAC AAA GAG) (SEQ ID NO: 14) and the same reverse primer as described in the Examples can be used to produce a PCR product that is approximately 750 base pairs in size and has an AatII restriction endonuclease site at the 5' end and an XbaI restriction endonuclease site at the 3' end. The product codes for amino acids 284-530 and includes the procollagen homology region (exons 6 and 7) and type 1 repeats. If inclusion of the TGF- β activating sequence (RFK) that is in the first type 1 repeat is found to reduce the antitumor activity, this sequence will be mutated to an inactive sequence (QFK, for example) using an oligonucleotidedirected mutagenesis kit (Amersham). The COMP/TSP-1P expression vector can be 15 constructed by cutting the PCR product with AatII and XbaI and cloning it into the COMP cDNA cut with the same enzymes. The protein can be expressed using the methods that have been described for COMP/TSP-1 and COMP/TSP-2.

Construction of COMP/Endostatin Expression Vectors

The strategy for making multimers of the TSP-1 and TSP-2 can be used to

20 make multimers of other anti-angiogenic proteins. For example, if the active region
of endostatin is prepared by PCR and cloned into the COMP cDNA, a pentameric
structure of endostatin can be made when this construct is expressed (O'Reilly M.D.,
et al., Cell 88:277-285, (1997)). In addition, if the COMP/TSP-1 and the
COMP/endostatin genes are expressed concurrently within the same cells, mixed

25 pentamers of COMP/TSP-1 and COMP/endostatin subunits are made. The mixed
multimer allows simultaneous treatment with the two reagents by delivery of a
single therapeutic. An additive or synergistic effect of the two agents may
significantly increase the efficacy of this reagent as compared to that of each reagent
alone. For example, combination therapy with angiostatin and endostatin has
eradicated tumors in mice (Boehm, T. et al., Nature 390:404-407, 1997).

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The cDNA for endostatin can be prepared by PCR of liver cDNA or from an isolated cDNA clone for collagen XVIII (GenBank accession no. L22548). The human endostatin cDNA can be produced by PCR with the forward primer GAT GAC GTC CAC AGC CAC CGC G (SEQ ID NO: 15) and the reverse primer GAT TCT AGA CTA CTT GGA GGC AGT CAT G (SEQ ID NO: 16). The resulting PCR product is approximately 560 base pairs and encodes amino acids 1 to 184 of human endostatin (Sasaki, T., et al., EMBO J., 17:4249-4256, 1998). The COMP/endostatin expression vector can be constructed by cutting the PCR product with AatII and XbaI, and cloning it into cDNA cut with the same enzymes. The protein can be expressed using the methods that have been described herein for COMP/TSP-1 and COMP/TSP-2. Angiostatin, as it was isolated from mice bearing Lewis lung carcinoma, includes the first four kringle domains of plasminogen (amino acids 98-440) (O'Reilly, M.S., et al., Cell 79:315-328, 1994). It should be noted that smaller constructs that contain fewer kringle domains should also be active based on published data (Griscelli, F., et al., Proc. Natl. Acad. Sci. USA 95:6367-6372, 1998). A 16,000 dalton fragment of prolactin and platelet factor 4 have also been reported to inhibit angiogenesis (Clapp, C. et al., Endocrinology 133:1292-1299, 1993; Gapta, S.K., et al., Proc. Natl. Acad. Sci. USA 92:7799-7803, 1995).

Also included in the inventions are compositions containing, as a biological ingredient, an anti-angiogenic chimeric protein, or a variant thereof to inhibit angiogenesis in mammalian tissues, and use of such compositions in the treatment of diseases and conditions characterized by, or associated with, angiogenic activity. Such methods can involve administration by oral, topical, injection, implantation, sustained release, or other delivery methods that bring one or more anti-angiogenic chimeric proteins in contact with cells whose growth is to be inhibited.

The present invention includes a method of treating an angiogenesis-mediated disease with a therapeutically effective amount of one or more anti-angiogenic chimeric proteins. Angiogenesis-mediated diseases can include, but are not limited to, cancers, solid tumors, tumor metastasis, benign tumors (e.g., hemangiomas, acoustic neuromas, neurofibromas, trachomas, and pyogenic

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granulomas), rheumatoid arthritis, psoriasis, ocular angiogenic diseases (e.g., diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis), Osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularization, telangiectasia, hemophiliac joints, angiofibroma, and wound granulation.

"Cancer" means neoplastic growth, hyperplastic or proliferative growth or a pathological state of abnormal cellular development and includes solid tumors, nonsolid tumors, and any abnormal cellular proliferation, such as that seen in leukemia. As used herein, "cancer" also means angiogenesis-dependent cancers and tumors, i.e., tumors that require for their growth (expansion in volume and/or mass) an increase in the number and density of the blood vessels supplying them with blood. "Regression" refers to the reduction of tumor mass and size. As used herein, the term "therapeutically effective amount" means the total amount of each active component of the composition or method that is sufficient to show a meaningful benefit to a treated human or other mammal, i.e., treatment, healing, prevention or amelioration of the relevant medical condition, or an increase in rate of treatment, healing, prevention or amelioration of such conditions. More specifically, for example, a therapeutically effective amount of an anti-angiogenic chimeric protein can cause a measurable reduction in the size or numbers of tumors, or in their rate of growth or multiplication, compared to untreated tumors. Other methods of assessing a "therapeutically effective amount," can include the result that blood vessel formation is measurably reduced in treated tissues compared to untreated tissues.

One or more anti-angiogenic chimeric proteins may be used in combination with other compositions and procedures for the treatment of diseases. For example, a tumor may be treated conventionally with surgery, radiation, chemotherapy, or immunotherapy, combined with anti-angiogenic chimeric proteins, and then anti-angiogenic chimeric proteins may be subsequently administered to the patient to extend the dormancy of micrometastases and to stabilize and inhibit the growth of any residual primary tumor.

The compositions may further contain other agents which either enhance the activity of the protein or compliment its activity or use in treatment, such as

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chemotherapeutic or radioactive agents. Such additional factors and/or agents may be included in the composition to produce a synergistic effect with protein of the invention, or to minimize side effects. Additionally, administration of the composition of the present invention may be administered concurrently with other therapies, e.g., administered in conjunction with a chemotherapy, immunotherapy or radiation therapy regimen.

The angiogenesis-modulating composition of the present invention may be a solid, liquid or aerosol and may be administered by any known route of administration. Examples of solid compositions include pills, creams, and implantable dosage units. The pills may be administered orally, the therapeutic creams may be administered topically. The implantable dosage unit may be administered locally, for example at a tumor site, or may be implanted for systemic release of the angiogenesis-modulating composition, for example subcutaneously. Examples of liquid composition include formulations adapted for injection subcutaneously, intravenously, intravenously, and formulations for topical and intraocular administration. Examples of aerosol formulation include inhaler formulation for administration to the lungs.

The anti-angiogenic chimeric proteins can be provided as isolated and substantially purified proteins in pharmaceutically acceptable formulations (including aqueous or nonaqueous carriers or solvents) using formulation methods known to those of ordinary skill in the art. These formulations can be administered by standard routes. In general, the combinations may be administered by the topical, transdermal, intraperitoneal, intracranial, intracerebroventricular, intracerebral, intravaginal, intrauterine, oral, rectal or parenteral (e.g., intravenous, intraspinal, subcutaneous or intramuscular) route. In addition, the anti-angiogenic chimeric proteins may be incorporated into biodegradable polymers allowing for sustained release of the compound, the polymers being implanted in the vicinity of where drug delivery is desired, for example, at the site of a tumor, or implanted so that the anti-angiogenic chimeric proteins is slowly released systemically. Osmotic minipumps may also be used to provide controlled delivery of high concentrations of anti-angiogenic chimeric proteins through cannulae to the site of interest, such as directly

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into a growth or into the vascular supply to that growth. The biodegradable polymers and their use are described, for example, in detail in Brem et al. (1991) (J. Neurosurg. 74:441-446), which is hereby incorporated by reference in its entirety.

As used herein, the terms "pharmaceutically acceptable," as it refers to compositions, carriers, diluents and reagents, represents that the materials are capable of administration to or upon a mammal with a minimum of undesirable physiological effects such as nausea, dizziness, gastric upset and the like. The preparation of a pharmacological composition that contains active ingredients dissolved or dispersed therein is well understood in the art and need not be limited based on formulation. Typically, such compositions are prepared as injectables either as liquid solutions or suspensions, however, solid forms suitable for solution, or suspensions, in liquid prior to use can also be prepared. The preparation can also be emulsified, for example, in liposomes.

The dosage of the anti-angiogenic chimeric proteins of the present invention will depend on the disease state or condition being treated and other clinical factors such as weight and condition of the human or animal and the route of administration of the compound. It is to be understood that the present invention has application for both human and veterinary use. The methods of the present invention contemplate single as well as multiple administrations, given either simultaneously or over an extended period of time.

The present invention also encompasses gene therapy whereby a polynucleotide encoding one or more anti-angiogenic chimeric proteins or one or more variants thereof, is introduced and regulated in a patient. Various methods of transferring or delivering DNA to cells for expression of the gene product protein, otherwise referred to as gene therapy, are disclosed in *Gene Transfer into Mammalian Somatic Cells in Vivo*, N. Yang (1992) *Crit. Rev. Biotechnol.* 12(4):335-356, which is hereby incorporated by reference. Gene therapy encompasses incorporation of DNA sequences into somatic cells or germ line cells for use in either ex vivo or in vivo therapy. Gene therapy can function to replace genes, augment normal or abnormal gene function, and to combat infectious diseases and other pathologies.

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Strategies for treating these medical problems with gene therapy include therapeutic strategies such as identifying the defective gene and then adding a functional gene to either replace the function of the defective gene or to augment a slightly functional gene; or prophylactic strategies, such as adding a gene for the product protein that will treat the condition or that will make the tissue or organ more susceptible to a treatment regimen. For example, a gene encoding an anti-angiogenic chimeric protein may be inserted into tumor cells of a patient and thus inhibit angiogenesis.

Gene transfer methods for gene therapy fall into three broad categories:

physical (e.g., electroporation, direct gene transfer and particle bombardment),
chemical (e.g., lipid-based carriers, or other non-viral vectors) and biological (e.g.,
virus-derived vector and receptor uptake). For example, non-viral vectors may be
used which include liposomes coated with DNA. Such liposome/DNA complexes
may be directly injected intravenously into the patient. It is believed that the

liposome/DNA complexes are concentrated in the liver where they deliver the DNA
to macrophages and Kupffer cells. These cells are long lived and thus provide long
term expression of the delivered DNA. Additionally, vectors or the "naked" DNA
of the gene may be directly injected into the desired organ, tissue or tumor for
targeted delivery of the therapeutic DNA.

In vivo gene transfer involves introducing the DNA into the cells of the patient when the cells are within the patient. Methods include using virally mediated gene transfer using a noninfectious virus to deliver the gene in the patient or injecting naked DNA into a site in the patient and the DNA is taken up by a percentage of cells in which the gene product protein is expressed. Additionally, the other methods described herein, such as use of a "gene gun," may be used for in vitro insertion of anti-angiogenic chimeric proteins DNA or anti-angiogenic chimeric proteins regulatory sequences.

Chemical methods of gene therapy may involve a lipid based compound, not necessarily a liposome, to transfer the DNA across the cell membrane. Lipofectins or cytofectins, lipid-based positive ions that bind to negatively charged DNA, make a complex that can cross the cell membrane and provide the DNA into the interior of

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the cell. Another chemical method uses receptor-based endocytosis, which involves binding a specific ligand to a cell surface receptor and enveloping and transporting it across the cell membrane. The ligand binds to the DNA and the whole complex is transported into the cell. The ligand gene complex is injected into the blood stream and then target cells that have the receptor will specifically bind the ligand and transport the ligand-DNA complex into the cell.

Many gene therapy methodologies employ viral vectors to insert genes into cells. For example, altered retrovirus vectors have been used in *ex vivo* methods to introduce genes into peripheral and tumor-infiltrating lymphocytes, hepatocytes, epidermal cells, myocytes, or other somatic cells. These altered cells are then introduced into the patient to provide the gene product from the inserted DNA.

Viral vectors have also been used to insert genes into cells using in vivo protocols. To direct the tissue-specific expression of foreign genes, cis-acting regulatory elements or promoters that are known to be tissue-specific can be used. Alternatively, this can be achieved using *in situ* delivery of DNA or viral vectors to specific anatomical sites *in vivo*. For example, gene transfer to blood vessels *in vivo* was achieved by implanting *in vitro* transduced endothelial cells in chosen sites on arterial walls. The virus infected surrounding cells which also expressed the gene product. A viral vector can be delivered directly to the *in vivo* site, by a catheter for example, thus allowing only certain areas to be infected by the virus, and providing long-term, site specific gene expression. *In vivo* gene transfer using retrovirus vectors has also been demonstrated in mammary tissue and hepatic tissue by injection of the altered virus into blood vessels leading to the organs.

Viral vectors that have been used for gene therapy protocols include but are not limited to, retroviruses, other RNA viruses such as poliovirus or Sindbis virus, adenovirus, adeno-associated virus, herpes viruses, SV40, vaccinia and other DNA viruses. Replication-defective murine retroviral vectors have been widely utilized gene transfer vectors.

Carrier mediated gene transfer *in vivo* can be used to transfect foreign DNA into cells. The carrier-DNA complex can be conveniently introduced into body fluids or the bloodstream and then site-specifically directed to the target organ or

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tissue in the body. Both liposomes and polycations, such as polylysine, lipofectins or cytofectins, can be used. Liposomes can be developed which are cell specific or organ specific and thus the foreign DNA carried by the liposome will be taken up by target cells. Injection of immunoliposomes that are targeted to a specific receptor on certain cells can be used as a convenient method of inserting the DNA into the cells bearing the receptor. Another carrier system that has been used is the asialoglycoprotein/polylysine conjugate system for carrying DNA to hepatocytes for in vivo gene transfer.

The gene therapy protocol for transfecting anti-angiogenic chimeric proteins into a patient may either be through integration of a gene encoding an antiangiogenic chimeric protein into the genome of the cells, into minichromosomes or as a separate replicating or non-replicating DNA construct in the cytoplasm or nucleoplasm of the cell. Anti-angiogenic chimeric proteins expression may continue for a long-period of time or may be reinjected periodically to maintain a desired level of the anti-angiogenic chimeric proteins protein in the cell, the tissue or organ or a determined blood level.

EXAMPLES

Example 1: Construction of COMP/TSP-1 and COMP/TSP-2

The chimeric expression vectors have been produced from three distinct cDNAs. The first is a clone for human cartilage oligomeric matrix protein (COMP) and was isolated from a λ gtll chondrocyte cDNA library (Doege, K.J, et al., J. Biol. Chem. 266:894-902 (1991)). This is an almost full-length clone for the COMP mRNA that only lacks a small region of the 5'-untranslated region. This clone (hCOMP-95) was used previously to determine the sequence of human COMP (GenBank Accession No. L32137; Genomics, 24:435-439 (1994)). 25

The second cDNA was produced using the polymerase chain reaction (PCR) with the human thrombospondin-1 (TSP-1) gene as the template. The TSP-1 clones were isolated from a human endothelial cell library (J. Cell Biol. 103:1635-1648

(1986)). The forward primer (GAT GAC GTC GAT GGT GGC TGG AGC CAC)
(SEQ ID NO: 17) and the reverse primer (GAT CTA GAT TGG ACA GTC CTG CTT G) (SEQ ID NO: 18) produce a PCR product that is approximately 354 basepairs in size and has an Aat II restriction endonuclease site at the 5' end and an
5 Xba I restriction endonuclease site at the 3' end. The PCR product encodes amino acids 417 to 530 and includes the second and third type 1 repeats of TSP-1 (see Figure 1 for the numbering of amino acids in TSP-1). The coding sequence for the first type 1 repeat was not included in the PCR product, by design, because it contains an RFK sequence that has been shown to activate TGF-β. This activity is not required to inhibit angiogenesis and it may produce unwanted secondary effects on numerous cell types. Vectors that include the first type 1 repeat can be constructed, using the same approach, if this region is found to enhance the antiangiogenic activity or other activities.

15 (catalog no. 936208 from Stratagene, LaJolla, CA) as the template. The forward primer (GAT GAC GTC GAG GAG GGC TGG TCT CCG) (SEQ ID NO: 19) and the reverse primer (GAT CTA GAC ACG GGG CAG CTC CTC TTG) (SEQ ID NO: 20) produced a PCR product that is approximately 520 base pairs in size and has an Aat II restriction endonuclease site at the 5' end and an Xba I restriction endonuclease site at the 5' end and an Xba I restriction of TSP-2 and, includes all three type 1 repeats of TSP-2 (see Figure 2 for numbering of amino acids in TSP-2). The sequence of the PCR primers was based on the human TSP-2 sequence in the GenBank database (Accession No. L12350). The sequences of the PCR products were determined to establish that mutations that affect the amino acid sequence had not been introduced during the PCR.

The COMP/TSP-1 and COMP-TSP-2 expression vectors were constructed by cutting the PCR products with Aat II and Xba I and subcloning them into the COMP cDNA vector [derived from Bluescript (Stratagene, La Jolla, CA)] cut with the same enzymes. The portion of COMP that was retained includes the signal sequence, the regions required for pentamerization and the first type 2 repeat (amino acids 1 to 128 on the enclosed sequence; Figure 3). Since there was an internal Aat

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II site in the TSP-2 PCR product, it had to be cloned into the vector in two steps. A 430 basepair Aat II/Xba I fragment of the TSP-2 PCR product was subcloned into the vector containing the portion of COMP as a first step. The resulting subclone was cut with Aat II, and a 90 base pair Aat II fragment of the PCR product was ligated into the expression vector. The final forms of the cDNAs were confirmed to have the predicted structure by nucleotide sequencing. They were then cut with Eco R1 and Xba I and ligated into the pcDNA 3.1 (Invitrogen; Carlsbad, CA) vector cut with the same enzymes. The DNA sequences of COMP/TSP-1 and COMP/TSP-2 are shown in Figures 4A and 4B and Figures 5A and 5B, respectively. The predicted molecular weights of the subunits of COMP/TSP-1 and COMP/TSP-2 should be approximately 24,200 and 30,000, respectively. The fully assembled COMP/TSP-1 and COMP/TSP-2 proteins should be 121,000 Da and 150,000 Da, respectively. The amino acid sequences of these proteins are shown in Figures 4A and 4B and Figures 5A and 5B, respectively.

15 Example 2: Production of Isolated COMP/TSP-1 and COMP/TSP-2

To express these chimeric proteins, the expression vectors can be transfected into human kidney 293 cells using the Lipofectin protocol (Gibco Laboratories). The cells can be selected with Zeocin and individual clones can be grown. The secretion of COMP/TSP-1 and COMP/TSP-2 can be monitored with western blotting using polyclonal antibodies to the region of COMP that is present in both expressed proteins. These antibodies have been produced by immunizing rabbits with a synthetically produced peptide, having an amino acid sequence derived from the N-terminal end of COMP, linked to a carrier protein. The amino acid sequence of the peptide is: SDLGPQMLRELQETN (SEQ ID NO: 21). A clone that expresses high levels of the protein can be grown in large volume flasks and in serum free media.

Example 3: Inhibition of Tumor Growth by COMP/TSP-1

A cDNA of thrombospondin-1 (TSP-1) containing the second and third type-1 repeats and the COMP assembly sequence (COMP/TSP-1) was produced by PCR

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using constructs derived as above as template, and was cloned into the expression vector pNeo (Invitrogen, Carlsbad, CA). Both the resulting COMP/TSP-1 construct and the unaltered vector alone were transfected into the human squamous carcinoma cell line A431 (Streit, M., et al., American Journal of Pathology 155:441-452, 1999), and positive clones were selected using Geneticin at a concentration of 800 µg/ml. The growth curves of positive clones were determined over an 8 day period. Clones of pNeo- and COMP/TSP-1 construct-transfected cells that had similar growth curves were selected to test the effect of the chimeric protein on tumor growth in nude mice. A total of five mice pre group were injected intradermally at the shoulders with 5 X 10⁶ cells per site, two sites per mouse. Every week the tumors were measured with calipers. At three weeks, the mice were sacrificed and the tumors were removed for further studies. As can be seen from Figure 7, expression of COMP/TSP-1 caused inhibition of the growth of the tumors in this model.

All references (e.g., journal articles, books, published patent applications and patents, etc.) cited herein are hereby incorporated by reference.

While this invention has been particularly shown and described with references to preferred embodiments thereof, it will be understood by those skilled in the art that various changes in form and details may be made therein without departing from the spirit and scope of the invention as defined by the appended claims.

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CLAIMS

What is claimed is:

- 1. A nucleic acid molecule encoding a chimeric protein comprising the second and third type 1 repeats of human TSP-1, but not the TGF- β activation region of human TSP-1.
- 2. A nucleic acid molecule encoding a chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the second and third type 1 repeats of human TSP-1.
- A nucleic acid molecule encoding a chimeric protein comprising the
 multimerization domain of human COMP, the first type 2 repeat of human
 COMP, and the second and third type 1 repeats of human TSP-1, but not the
 TGF-β activation region of human TSP-1.
 - 4. A nucleic acid molecule encoding a chimeric protein comprising the multimerization domain of human COMP, the procollagen homology region of TSP-1, and the first, second, and third type 1 repeats of human TSP-1.
 - 5. A nucleic acid molecule encoding a chimeric protein comprising the multimerization domain of human COMP, the procollagen homology region of TSP-1, and the first, second, and third type 1 repeats of human TSP-1, but not the TGF-β activation region of human TSP-1.
- A nucleic acid molecule encoding a chimeric protein comprising the multimerization domain of human COMP and a portion of human endostatin, wherein the chimeric protein has anti-angiogenic activity.

- 7. A nucleic acid molecule encoding a chimeric protein comprising the multimerization domain of human COMP and a portion of human angiostatin, wherein the chimeric protein has anti-angiogenic activity.
- 8. A nucleic acid molecule encoding a chimeric protein comprising the
 5 multimerization domain of human COMP and a portion of human prolactin,
 wherein the chimeric protein has anti-angiogenic activity.
 - 9. A nucleic acid molecule encoding a chimeric protein comprising the multimerization domain of human COMP and a portion of human platelet factor 4, wherein the chimeric protein has anti-angiogenic activity.
- 10 10. A nucleic acid molecule encoding a chimeric protein comprising the multimerization domain of human COMP, the procollagen homology region, and the first, second, and third type 1 repeats of human TSP-1.
 - 11. A nucleic acid molecule encoding a protein having the amino acid sequence SEQ ID NO: 5.
- 15 12. A vector comprising nucleic acid encoding a chimeric protein comprising the second and third type 1 repeats of human TSP-1 but not the TGF-β activation region of human TSP-1.
 - 13. A host cell comprising the vector of Claim 12.
- 14. A vector comprising nucleic acid encoding a chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the second and third type 1 repeats of human TSP-1.
 - 15. A host cell comprising the vector of Claim 14.

- 16. A method for producing a chimeric protein which comprises the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the second and third type 1 repeats of human TSP-1, said method comprising maintaining the host cell of Claim 15 under conditions suitable for expression of said nucleic acid, whereby said protein is produced.
- 17. The method of Claim 16 further comprising isolating the chimeric protein.
- A vector comprising nucleic acid encoding a chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human
 COMP, and the second and third type 1 repeats of human TSP-1, but not the TGF-β activation region of human TSP-1.
 - 19. A host cell comprising the vector of Claim 18.
- A method for producing a chimeric protein which comprises the multimerization domain of human COMP, the first type 2 repeat of human
 COMP, and the second and third type 1 repeats of human TSP-1, but not the TGF-β activation region of human TSP-1, said method comprising maintaining the host cell of Claim 19 under conditions suitable for expression of said nucleic acid, whereby said protein is produced.
 - 21. The method of Claim 20 further comprising isolating the chimeric protein.
- 20 22. A vector comprising nucleic acid encoding a chimeric protein comprising the multimerization domain of human COMP, the procollagen homology region, and the first, second, and third type 1 repeats of human TSP-1.
 - 23. A vector comprising nucleic acid encoding a protein having the amino acid sequence SEQ ID NO: 5.

- 24. A host cell comprising the vector of Claim 23.
- 25. A chimeric protein comprising the second and third type 1 repeat of human TSP-1, but not the TGF- β activation region of human TSP-1.
- A chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the second and third type 1 repeats of human TSP-1.
 - 27. A chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the second and third type 1 repeats of human TSP-1, but not the TGF-β activation region of human TSP-1.
 - A chimeric protein comprising the multimerization domain of human COMP, the procollagen homology region of TSP-1, and the first, second, and third type 1 repeats of human TSP-1.
- A chimeric protein comprising the multimerization domain of human COMP and a portion of human endostatin, wherein the chimeric protein has antiangiogenic activity.
 - 30. A chimeric protein comprising the multimerization domain of human COMP and a portion of human angiostatin, wherein the chimeric protein has antiangiogenic activity.
- 20 31. A chimeric protein comprising the multimerization domain of human COMP and a portion of human prolactin, wherein the chimeric protein has anti-angiogenic activity.

- 32. A chimeric protein comprising the multimerization domain of human COMP and a portion of human platelet factor 4, wherein the chimeric protein has anti-angiogenic activity.
- 33. A protein having the amino acid sequence SEQ ID NO: 5.
- 5 34. An isolated nucleic acid molecule encoding a chimeric protein comprising the three type 1 repeats of human TSP-2.
 - 35. A vector comprising nucleic acid encoding a chimeric protein comprising the three type 1 repeats of human TSP-2.
 - 36. A host cell comprising the vector of Claim 35.
- 10 37. A method for producing a chimeric protein which comprises the three type 1 repeats of human TSP-2, said method comprising maintaining the host cell of Claim 36 under conditions suitable for expression of said nucleic acid, whereby said protein is produced.
 - 38. The method of Claim 37 further comprising isolating the chimeric protein.
- A nucleic acid molecule encoding a chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the three type 1 repeats of human TSP-2.
 - 40. A vector comprising isolated nucleic acid encoding a chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the three type 1 repeats of human TSP-2.
 - 41. A host cell comprising the vector of Claim 40.

- 42. A method for producing a chimeric protein which comprises the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the three type 1 repeats of human TSP-2, said method comprising maintaining the host cell of Claim 41 under conditions suitable for expression of said nucleic acid, whereby said protein is produced.
- 43. The method of Claim 42 further comprising isolating the chimeric protein.
- 44. A nucleic acid molecule encoding a protein having the amino acid sequence SEQ ID NO: 7.
- 45. A vector comprising nucleic acid encoding a protein having the amino acid sequence SEQ ID NO: 7.
 - 46. A host cell comprising the vector of Claim 45.
 - 47. A chimeric protein comprising the three type 1 repeats of human TSP-2.
 - 48. A chimeric protein comprising the procollagen homology region of TSP-2 and the three type 1 repeats of human TSP-2.
- A chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the three type 1 repeats of human TSP-2.
 - 50. A protein having the amino acid sequence SEQ ID NO: 7.
- A method for inhibiting angiogenesis in a human or other mammal, the
 method comprising administering to the human or other mammal a
 therapeutically effective amount of an anti-angiogenic chimeric protein.

- 52. The method of Claim 51 wherein the anti-angiogenic chimeric protein is selected from the group consisting of:
 - a) a chimeric protein comprising the second and third type 1 repeats of human TSP-1;
- b) a chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the second and third type 1 repeats of human TSP-1;
 - c) a chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the second and third type 1 repeats of human TSP-1, but not the TGF-β activation region of human TSP-1;
 - d) a chimeric protein comprising the multimerization domain of human COMP, the procollagen region, and the first, second, and third type 1 repeats of human TSP-1; and
- a chimeric protein comprising the three type 1 repeats of human TSP-2; and (6) a chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the three type 1 repeats of human TSP-2.
- 53. The method of Claim 51 wherein the anti-angiogenic protein is administered locally at the site of one or more growths.

	human	thrombospondin-1
NH ₂	1 61 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	NRIPESGGDNSVFDIFELTGAARKGSGRRLVKGPDPSSPAPRIEDANLIPPVPDDKFQDL VDAVRTEKGFLLLASLRQHKKTRGTLLALERKDHSGQVFSVVSNGKAGTLDLSLTVQGKQ HVVSVEBALLATGQWKSITLFVQEDRAQLYIDCEKHENARLDVPIQSVFTRDLASIARLR IAKGGVNDNFQGVLQNVRFVFGTTPRDILRNKGCSSSTSVLLTLDNNVV®GSSPAIRTNY
	241	IGHKTKDLQAI¢GIS¢DRLSSH
<u>.</u>	312	VLELRGLRTIVTTLQDSIRKVTEENKELANBLRRPPLCYHNGVQYRNNE BWTVDS.CTECHCQNSVTICKKVSCPIMPCS®ATVPDGECCPRCWPSDSA
type 1	361 417 474	DDGWSPWSEWTSCSTSCGNGIQQRGRSCDSLNNRCRGSSVQTRTCHIQRCDKRFKQ DGGWSRWSPWSSGS <u>VTCB</u> DGVITRIRLCNSPSPQMNGKPCBGRARETKACKKDACPI NGGWGPWSPWDICS <u>VTCG</u> GGVQKRSRLCN(MPTPQFGGKDCVGDVTENQICNKQDCPI
type 2	531 572 630	DGCLSNPCFAGVKCTSYPDGSWKCGACPPGYSGNGIQCTDV DECKEVPDACFNHNGEHRCENTDPGYNCLPCPPRFTGSQPFGQGVBHATANKQVCKPR NPCTDGTHDCNKNAKCNYLGHYSDPMYRC-RCKPGYAGNGIICGE
	674	DTDLDGWPNENLVCVA(MATTHCKK
type 3	698 734 757 793 793 816	DNCPNLPNSGQBDYDKDGIGDACDDDDDNDKIPDDR DNCPFHYNPAQYDYDRDDVGDRC DNCPYHHNPDQADTDNNGEGDACAADIDGDGILNBR DNCQYVYNVDQRDTDMDGVGDQC DNCPLEHNPDQLDSDSDRIGDTCDNNQDIDEDGHQNNL DNCPLYPNANQADHDKDGKGDACDHDDDNDGIPDDK
СООН	926 986 1046	DICPENVDISKTDPRRFQHIPLDPRGTSQNDPNWVVRHQGKELVQTVNCDPGLAVGYDBP NAVDFSGTPPIHTRRDDDYAGFVFGYQSSR <u>RYVVMMK</u> QVTQSYWDTMPTRAQGYSGLSV KVV(W)STTGPGRHLRNALWHTGNTPGQVRTLWHDPRHIGWKDFTAYRWRLSHRPKTGF <u>IRV</u> VHYRGKKIMADSGPIYDKTYAGGRLGLFVFSQRHVFFSDLKYECRDP

FIG

ц	human thr	thrombospondin-2
NH ₂	1 NVWR 61 DYIP 121 ADTI	VWRLVLLALHVHFSTQAGUQDKDTTFDLFSISNINRKTIGARQFRGPDPGVPAYRFVRF YIPPVNADDLSKITKIHRQKEGFFLTAQLKQDGKSRGTLLALEGPGLSQRQFEIVSNGP DTLDLTYWIDGTRHVVSLEDVGLADSQHKØVTVQVAGETYSLHVGCDLIGPVALDEPFY HLQARKSRHYVAKGBARBSHPRGLLQNVHLVFENSVEDILSKKGCQQGQGAEINAISEN
	241 TETI	ETLRLGPHVTTEYVGPSSKRRPEVCERSCEELGNM
ū.	277 VQEI 332 TWV	QELSGLHVLVNQLSENLKRVSNDNQFLWELIGGPPKTRØMSACWQDGRFFAE@B RVVDSCTTCTCKKFKTICHQITCPPATCASPSFVEGECCPSCLHSVDG
type 1	381 KEGW 437 DGGW 494 DGRW	GWSPWAEWTQCS <u>VYCG</u> SGTQQRGRSCDVTBNTCLGPSIQTRACSLSKCDTRIRQ GWSHWSPWSSCS <u>VYCG</u> VG(B)ITRIRLCNSPVPQMGGKNCKGSGRKTKACQGAPCPI RWSPWSACTVTCAGGIRERTRVCNSPRPQYGGKACVGDVQERQKCNKRSCPV
type 2	551 DGC 592 DBC 650 NPC	GCLSNPCFPGAQCSSPPDGSWSCGFCPVGFLG
	694 DSD	SDLDGWPHINLVCAT®ATYHCIK
type 3	718 DNC 748 DNC 777 DNC 813 DNC 836 DNC 874 DNC	DNCPHLPNSGQRDFDKDGIGDACDDDDDNDGVTDRK DNCQLLFNPRQADYDKDBVGDRC BNCPYVHNPAQIDTDNNGBGDACSVDIDGDDVFNER BNCPYVYNTDQRDTDGDGVGDHC DNCPLVHNPDQTDVDNDLVGDQCDNNEDIDDDGHQNNQ DNCPLYHNPDQEDDGGGDACDPDDDNDGVPDDR
СООН	946 DVC 1006 GSV 1066 KV	DVCPENHAISETDERHFQMVPLDPKGTTQIDPNHVIRHQGKELVQTANSDPGIAVGFDBF GSVDFSGTFYVNTDRDDDYAGFVFGYQSSSR <u>FYVVHMR</u> QVTQTYWBDQPTRAYGYSGVSL KVVWBSTGTGEHLRHALHHTGNTPGQVRTLHHDPRNIGHRDYTAYRHHLTHRPKTGYIRV LVHEGKQVHADSGPIYDQTYAGGRLGLFVFSQEHVYPBDLKYECRDI

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VPDTA CV LLLTLAALGASGQGQSPLGSDLGPQMLRELQETNA
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human COMP

LODVRDWLRQQVREIT 61 FLKNTVMECDACGMQQBVRTGLPSVRPL

Hype 2 182 NECETGOHN-CVPNSVCINTRGSFQ-CGPCQPGFVG-------DQASGCQRGAQ 129 NECNAHP---CFPRVRCI@TSPGFR-CEACPPGYSGPTHQGVGLAFAKANKQVCTDI ----NGILCGR 227. RFCPDGSPSECHEHADCVLERDGSRSCV-CRVGWAG----89 LHCAPGF---CPPGVACIQTESGGR-CGPCPAGFTG--

269 DTDLDGFPDEKLRCPEPQCRK

290 DNCVTVPNSGQEDVDRDGIGDACD--PDADGDGVPNEK DNCPLVRNPDQRNTDEDKWGDAC 326

DNCPQKSNPDQADVDHDFVGDACDSDQDQDGHQDSR DNCPTVPNSAQEDSDHDGQGDACD--DDDDNDGVPDSR DNCPRVPNSDQKDSDGDGIGDAC 408 446

DNCRLVPNPGQEDADRDGVGDVCQ--DDFDADKVVDKI

518 DYCPENARYTLTDFRAFQTVVLDPEGDAQIDPNWVLNQGREIVQTHNSDPGLAVGYTAF NG V D F E G T F H V N T V T D D D Y A G F I P G Y Q D S S S F Y V V M N K Q H E Q T Y W Q A N P F R A V A E P G I Q L KAVKSSTGPGRQLRNALWHTGDTESQVRLLWKDPRHVGNKDKKSYRWFLQHRPQVGYIRV REYRGPELVADSNVVLDTTMRGGRLGVFCFSQBNITWANLRYRCHDTIPEDYETHQLRQA COOH 578

F16. 3

CAGO	ACCC	'AG C	TCC	CGCC	CA CO	GCC	ATG	GTC	ccc	GAC	ACC	GCC	TGC	GTT	CTT	52
														Val		
CTG Leu 10	CTC Leu	ACC Thr	CTG Leu	GCT Ala	GCC Ala 15	CTC Leu	GJ Y GGC	GCG Ala	TCC Ser	GGA Gly 20	CAG Gln	Gly	CAG Gln	AGC Ser	CCG Pro 25	100
TTG Leu	Gly	TCA Ser	GAC Asp	CTG Leu 30	GGC Gly	CCG Pro	CAG Gln	ATG Met	CTT Leu 35	Arg	GAA Glu	CTG Leu	CAG Gln	GAA Glu 40	ACC Thr	148
AAC Asn	GCG Ala	GCG Ala	CTG Leu 45	CAG Gln	GAC Asp	GTG Val	cgg Arg	GAC Asp 50	TGG Trp	CTG Leu	Arg	CAG Gln	ÇAG Gln 55	GTC Val	AGG Arg	196
GAG Glu	ATC Ile	ACG Thr 60	TTC Phe	CTG Leu	AAA Lys	AAC Asn	ACG Thr 65	GTG Val	ATG Met	GAG Glu	TGT Cys	GAÇ Asp 70	GCG Ala	TGC Cys	GGG	244
ATG Met	CAG Gln 75	CAG Gln	TCA Ser	GTA Val	CGC Arg	ACC Thr 80	GGC	CTA Leu	CCC Pro	AGC Ser	GTG Val 85	CGG	CCC Pro	CTG	CTC Leu	292
CAC His 90	TGC Cys	GCG Ala	CCC	GGC	TTC Phe 95	TGC. Cys	TTC Phe	CCC Pro	GGC	GTG Val 100	GCC Ala	TGC Cys	ATC Ile	CAG Gln	ACG Thr 105	340
GAG Glu	AGC Ser	GGC	GGC	CGC Arg 110	TGC Cys	GGC	CCC Pro	TGC Cys	CCC Pro 115	GCG Ala	GGC Gly	TTC Phe	ACĠ Thr	GGC Gly 120	AAC Asn	388

FIG 4A

5/9

ggc ely	TCG Ser	CAC His	TGC Cys 125	ACC Thr	GAC Asp	GTC Val	GAT Asp	GGT Gly 130	GGC Gly	TGG Trp	AGC Ser	CAC His	TGG Trp 135	TCC	CCG Pro	436
TGG Trp	TCA Ser	TCT Ser 140	TGT Cys	TCT Ser	GTG Val	ACA Thr	TGT Cys 145	GGT Gly	GAT Asp	GGT Gly	GTG Val	ATC Ile 150	ACA Thr	AGG Arg	ATC Ile	484
CGG Arg	CTC Leu 155	TGC Cys	AAC Asn	TCT Ser	CCC Pro	AGC Ser 160	CCC Pro	CAG Gln	ATG Met	AAC Asn	GGG Gly 165	AAA Lys	CCC Pro	TGT	GAA Glu	532
GGC Gly 170	GAA Glu	GCG Ala	CGG Arg	GAG Glu	ACC Thr 175	AAA Lys	GCC	TGC Cys	AAG Lys	AAA Lys 180	GAC Asp	GCC Ala	TGC	CCC Pro	ATC Ile 185	580
AAT Asn	GGA Gly	GGC Gly	TGG Trp	GGT Gly 190	CCT Pro	TGG Trp	TCA Ser	CCA Pro	TGG Trp 195	GAC Asp	ATC Ile	TGT Cys	TCT Ser	GTC Val 200	ACC Thr	628
TGT Cys	GGA Gly	GGA Gly	GGG Gly 205	GTA Val	CAG Gln	AAA Lys	CGT Arg	AGT Ser 210	CGT Arg	CTC Leu	TGC Cys	AAC Asn	AAC Asn 215	CCC Pro	ACA Thr	676
						Asp						ACA Thr 230				724
		AAC Asn								A						755

FIG. 4B

CAGC	'ACCC	AG C	TCCC	CGCC	CA CC	GCC	ATG Met 1	GTC Val	CCC Pro	GAC Asp	ACC Thr 5	GCC Ala	CYB	GTT Val	CTT Leu	52
CTG Leu 10	CTC Leu	ACC Thr	CTG Leu	GCT Ala	GCC Ala 15	CTC Leu	GGC Gly	GČG Ala	TCC Ser	GGA Gly 20	CAG Gln	GGC	CAG Gln	AGC Ser	CCG Pro 25	100
TTG Leu	GGC Gly	TCA Ser	GAC Asp	CTG Leu 30	GGC Gly	CCG Pro	CAG Gln	ATG Met	CTT Leu 35	CGG Arg	GAA Glu	CTG Leu	CAG Gln	GAA Glu 40	ACC Thr	148
AAC Asn	GCG Ala	GCG Ala	CTG Leu 45	CAG Gln	GAC Asp	GTG Val	CGG Arg	GAC Asp 50	TGG Trp	CTG Leu	cgg Arg	CAG Gln	CAG Gln 55	GTC Val	AGG Arg	196
GAG Glu	ATC Ile	ACG Thr 60	TTC Phe	CTG Leu	AAA Lys	AAC Asn	ACG Thr 65	GTG Val	ATG Met	GAG Glu	TGT Cys	GAC Asp 70	GCG Ala	TGC Cys	GGG	244
Met	Gln 75	Gln	Ser	Val	CGC Arg	Thr 80	Gly	Leu	Pro	ser	85	Arg				292
CAC His 90	TGC Cys	GCG Ala	CCC Pro	GGC Gly	TTC Phe 95	TGC Cys	TTC Phe	CCC Pro	GGC Gly	GTG Val 100	GCC Ala	TGC	ATC Ile	CAG Gln	ACG Thr 105	340
GAG Glu	AGC Ser	GGC Gly	GGC Gly	CGC Arg 110	TGC Cys	GGC Gly	CCC	TGC	CCC Pro 115	GCG Ala	GGC	TTC Phe	ACG Thr	GGC Gly 120	AAC Asn	388
GGC Gly	TCG Ser	CAC His	TGC Cys 125	ACC Thr	GAC Asp	GTC Val	GAG Glu	GAG Glu 130	GGC	TGG	TCT Ser	CCG Pro	TGG Trp 135	GCA Ala	GAG Glu	436
TGG	ACC Thr	CAG Gln 140	Сув	TCC Ser	GTG Val	ACG Thr	TGT Cys 145	GIA	TCT Ser	ggg ġly	ACC Thr	CAG Gln 150	<u> </u>	AGA Arg	GGC Gly	484

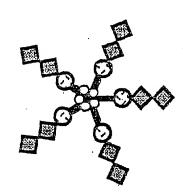
FIG. 5A

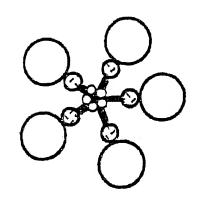
	Cys								CAG Gln		532
Arg	GCT Ala										580
	TGG Trp										628
	GGC Gly										676
	GGG Gly 220								GCC Ala	·	724
	GGC Gly										772
	.TCG Ser							Glu			820
	GTC Val		_		Pro						868
	GAT Asp			Gln							916
TCT Ser											925

F1G. 5B

COMP/TSP-1

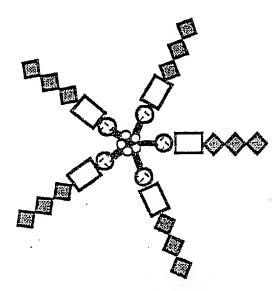
COMP/ENDOSTATIN

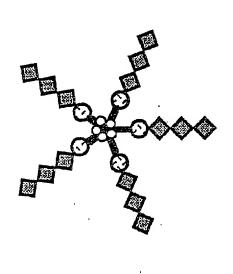




COMP/TSP-1P

COMP/TSP-2







pentamerization domain of human COMP



type 2 repeat of human COMP



second and third type 1 repeats of TSP-1



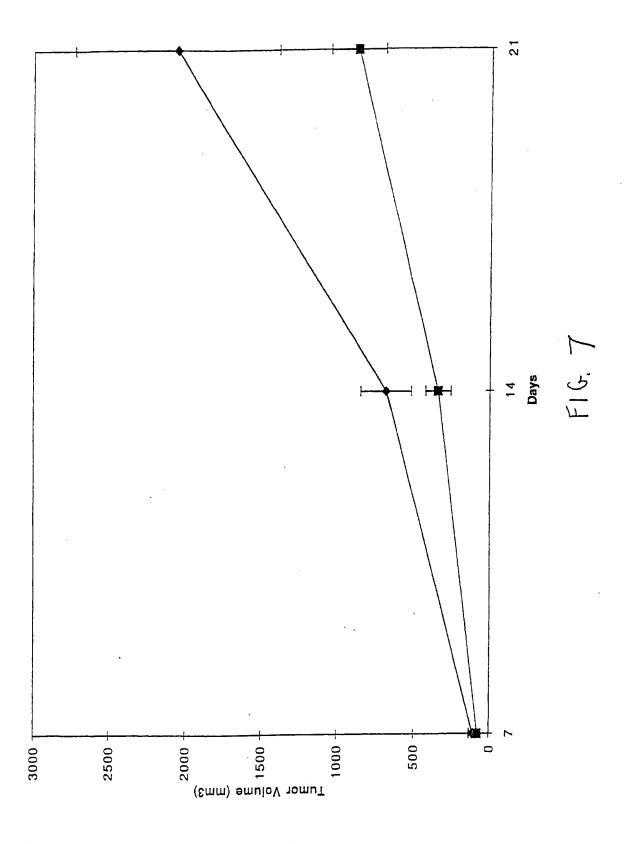
all three type1 repeats of TSP-1 or -2



procellagen homology region



endostatin



(19) World Intellectual Property Organization International Bureau



(43) International Publication Date 3 August 2000 (03.08.2000)

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- (21) International Application Number: PCT/US00/02482
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- (25) Filing Language:

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- (71) Applicant (for all designated States except US): BETH ISRAEL DEACONESS MEDICAL CENTER [US/US]; 330 Brookline Avenue, Boston, MA 02215 (US).
- (72) Inventor; and
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- (74) Agents: HOGLE, Doreen, M. et al.; Hamilton, Brook, Smith & Reynolds, P.C., Two Militia Drive, Lexington, MA 02421 (US).

- (81) Designated States (national): AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW.
- (84) Designated States (regional): ARIPO patent (GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).

Published:

- With international search report.
- (88) Date of publication of the international search report: 15 February 2001

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

(54) Title: COMP/TSP-1, COMP/TSP-2 AND OTHER TSP CHIMERIC PROTEINS

(57) Abstract: Tumors attract blood vessels in order to grow by a process called angiogenesis. The relative quantity of stimulators and inhibitors is an important determining factor for the initiation of angiogenesis. Thrombospondins-1 and -2 are adhesive glycoproteine that have the ability to inhibit aniogenesis. This inhibiting activity has been mapped to the type 1 repeats of TSP-1 and TSP-2. The invention includes chimeric proteins that contain anti-angiogenic portions of TSP-1, TSP-2, endostatin, angiostatin, platelet factor 4, or prolactin, linked to a portion of the N-terminal region of human cartilage oligomeric matrix protein (COMP) that allows formation of pentamers. Also described herein are the nucleic acid molecules, vectors, and host cells for expressing and producing these chimeric proteins. Further embodiments of the invention include methods to treat humans or other mammals with anti-angiogenic proteins to reduce tumor size or rate of growth. Since the type 1 repeat region of TSP-1 and TSP-2 reportedly inhibits HIV infection, chimeric proteins comprising these repeats may also be used for this purpose, as well as to inhibit angiogenesis.

Intern ial Application No PCT/US 00/02482

A. CLASSIFICATION OF SUBJECT MATTER IPC 7 C12N15/52 C07K A61K38/17 C07K19/00 C07K14/78 According to International Patent Classification (IPC) or to both national classification and IPC B. FIELDS SEARCHED Minimum documentation searched (classification system followed by classification symbols) IPC 7 CO7K C12N Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched Electronic data base consulted during the international search (name of data base and, where practical, search terms used) EPO-Internal, CHEM ABS Data, BIOSIS, MEDLINE, SCISEARCH, BIOTECHNOLOGY ABS, WPI Data C. DOCUMENTS CONSIDERED TO BE RELEVANT Relevant to claim No. Citation of document, with indication, where appropriate, of the relevant passages Category ' PANETTI TS ET AL: "Interaction of 1,12,13, χ 25 recombinant procollagen and properdin modules of thrombospondin-1 with heparin and fibrinogen/fibrin." JOURNAL OF BIOLOGICAL CHEMISTRY, JAN 1 1999, 274 (1) P430-7, XP002140107 UNITED STATES 2-5,10, Α abstract; figure 2B 11, 14-24, 26-28 Patent family members are listed in annex. Further documents are listed in the continuation of box C. χ Χ Special categories of cited documents: *T* later document published after the international filing date or priority date and not in conflict with the application but "A" document defining the general state of the art which is not considered to be of particular relevance clied to understand the principle or theory underlying the Invention "E" earlier document but published on or after the international *X* document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone "L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified) "Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents with the property to a proposition. *O" document referring to an oral disclosure, use, exhibition or ments, such combination being obvious to a person skilled other means in the art. *P* document published prior to the international filing date but later than the priority date claimed "&" document member of the same patent family Date of malling of the international search report Date of the actual completion of the international search 1 8, 10, 00 21 September 2000 Authorized officer Name and malling address of the ISA European Patent Office, P.B. 5818 Patentiaan 2 NL = 2280 HV Rijswijk Tel. (+31=70) 340=2040, Tx. 31 651 epo nl, Fax: (+31=70) 340=3016 Gurdjian, D

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Intern Aal Application No PCT/US 00/02482

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·	ation) DOCUMENTS CONSIDERED TO BE RELEVANT	
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Inte. utional application No. PCT/US 00/02482

Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)
This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:
Claims Nos.: because they relate to subject matter not required to be searched by this Authority, namely:
Although claims 51-53 are directed to a method of treatment of the human/animal body, the search has been carried out and based on the alleged effects of the compound/composition.
Claims Nos.: because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:
3. Claims Nos.:
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).
Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)
This International Searching Authority found multiple inventions in this international application, as follows:
see additional sheet
As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.
2. As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:
4. No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:
Remark on Protest The additional search fees were accompanied by the applicant's protest. X No protest accompanied the payment of additional search fees.

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

This International Searching Authority found multiple (groups of) inventions in this international application, as follows:

1. Claims: 51-53 partially and 1-5,10-28 ,33

TSP-1 chimeric protein constructs, corresponding nucleic acids ,vectors ,host cells , method of producing chimeric proteins

2. Claims: 51-53 partially and 34-50

TSP-2 chimeric protein constructs, corresponding nucleic acids ,vectors ,host cells ,method of producing chimeric proteins

3. Claims: 51-53 partially and 6-9, 29-32

COMP chimeric protein constructs, not provided in the first and second invention of present application, corresponding nucleic acids, vectors, host cells, method of producing chimeric proteins

Information on patent family members

Intern nal Application No PCT/US 00/02482

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CORRECTED VERSION

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(22) International Filing Date: 1 February 2000 (01.02.2000)

(25) Filing Language: English

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(71) Applicant (for all designated States except US): BETH ISRAEL DEACONESS MEDICAL CENTER [US/US]; 330 Brookline Avenue, Boston, MA 02215 (US).

(72) Inventor; and

(75) Inventor/Applicant (for US only): LAWLER, John, W. [US/US]; 6 Gale Road, Swampscott, MA 01907 (US).

(74) Agents: HOGLE, Doreen, M. et al.: Hamilton. Brook. Smith & Reynolds, P.C., Two Militia Drive, Lexington, MA 02421 (US).

(81) Designated States (national): AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK,

DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW.

(84) Designated States (regional): ARIPO patent (GH. GM. KE, LS, MW, SD, SL, SZ, TZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE). OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).

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(15) Information about Correction: see PCT Gazette No. 07/2002 of 14 February 2002, Section II

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

(54) Title: COMP/TSP-1, COMP/TSP-2 AND OTHER TSP CHIMERIC PROTEINS

(57) Abstract: Tumors attract blood vessels in order to grow by a process called angiogenesis. The relative quantity of stimulators and inhibitors is an important determining factor for the initiation of angiogenesis. Thrombospondins-1 and -2 are adhesive glycoproteine that have the ability to inhibit aniogenesis. This inhibiting activity has been mapped to the type 1 repeats of TSP-1 and TSP-2. The invention includes chimeric proteins that contain anti-angiogenic portions of TSP-1, TSP-2, endostatin, angiostatin, platelet factor 4, or prolactin, linked to a portion of the N-terminal region of human cartilage oligomeric matrix protein (COMP) that allows formation of pentamers. Also described herein are the nucleic acid molecules, vectors, and host cells for expressing and producing these chimeric proteins. Further embodiments of the invention include methods to treat humans or other mammals with anti-angiogenic proteins to reduce tumor size or rate of growth. Since the type 1 repeat region of TSP-1 and TSP-2 reportedly inhibits HIV infection, chimeric proteins comprising these repeats may also be used for this purpose, as well as to inhibit angiogenesis.



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COMP/TSP-1, COMP/TSP-2 AND OTHER CHIMERIC PROTEINS

RELATED APPLICATIONS

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This application claims the benefit of U.S. Provisional Application No. 60/118,053 filed February 1, 1999, the entire teachings of which are incorporated 5 herein by reference.

BACKGROUND OF THE INVENTION

Thrombospondins are a family of calcium-binding multifunctional glycoproteins that are secreted by various cell types and are developmentally regulated components of the extracellular matrix (Bornstein, P., FASEB J., 6:3290-10 3299, 1992; Bornstein, P., J. Cell Biol., 130:503-506, 1995). Among their functions are modulating cell attachment, migration and proliferation.

One member of this family, cartilage oligomeric matrix protein (COMP) is a pentamer in which multimerization appears to be directed by α -helical segments situated (in the amino acid sequence) either before or after the cysteine residues that 15 form the interchain disulfide bonds. COMP has been purified (Prochownik, E.V. et al., J. Cell Biol. 109:843-852 (1989)). Individuals affected with pseudoachondroplasia, who have considerably shortened stature as a result of premature cessation of bone growth, have been shown to have mutations in exon 17B of the COMP protein (Nature Genetics 10:325-329 (1995)).

In vitro assays have shown that platelet thrombospondin-1 is involved in thrombosis, fibrinolysis, wound healing, inflammation, tumor cell metastasis and angiogenesis. The major form of thrombospondin secreted by platelets and endothelial cells is TSP-1. Thrombospondin-1 (TSP-1) is an angiogenesis inhibitor that decreases tumor growth. Thrombospondin- 2 (TSP-2) is a related glycoprotein 25 of similar structure and properties.

The thrombospondin type 1 repeats (TSRs; also "repeat regions" herein) have been shown to inhibit angiogenesis and HIV infection. However, other portions of the proteins have been shown to have a positive effect on endothelial cell

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growth. Thromobospondin-1 and -2 are similar in terms of their molecular architecture. Thrombospondin-land thrombospondin-2 each have three copies of the TSR. TSP-1 and TSP-2 are trimeric molecules. Thus, each fully assembled protein contains nine TSRs.

Whereas TSP-1 and TSP-2 are antiangiogenic, these proteins contain other domains that have additional activities that diminish the antiangiogenic activity. The isolated TSRs are more potent inhibitors of angiogenesis than the native molecules.

The ingrowth of new capillary networks into developing tumors is essential for the progression of cancer. Thus, the development of pharmaceuticals that inhibit 10 the process of angiogenesis is an important therapeutic goal. As pointed out in a review by Folkman (Folkman, J., Proc. Natl. Acad. Sci. USA 95: 9064-9066, 1998), antiangiogenic therapy has little toxicity, does not require the therapeutic agent to enter tumor cells or cross the blood-brain barrier, controls tumor growth independently of growth of tumor cell heterogeneity, and does not induce drug 15 resistance.

SUMMARY OF THE INVENTION

The invention includes chimeric proteins comprising: (1) a chimeric protein comprising the second and third type 1 repeats of human TSP-1, and which may also 20 comprise the procollagen homology region of TSP-1; (2) a chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the second and third type 1 repeats of human TSP-1; (3) a chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the second and third type 1 repeats of human TSP-1, but not the TGF-β activation region of human TSP-1; (4) a chimeric protein comprising the multimerization domain of human COMP, the procollagen region, and the first, second, and third type 1 repeats of human TSP-1; (5) a chimeric protein comprising the three type 1 repeats of human TSP-2, and which may also comprise the procollagen homology region of TSP-2; (6) a chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP,

and the three type 1 repeats of human TSP-2; and (7) variants of any of the above having anti-angiogenic activity. The invention further includes isolated nucleic acids encoding any of the above chimeric proteins, vectors comprising these nucleic acids, and host cells comprising any of said vectors. The chimeric proteins can be produced in host cells and used in methods for the treatment of a disease or medical condition characterized by abnormal or undesirable proliferation of blood vessels, such as that occurring in tumor growth.

BRIEF DESCRIPTION OF THE DRAWINGS

Figure 1 is a representation of the amino acid sequence of human TSP-1 (SEQ ID NO: 1). The type 1 repeats of TSP-1 are, as illustrated here, 1) amino acids 361-416; 2) amino acids 417-473; and 3) amino acids 474-530.

Figure 2 is a representation of the amino acid sequence of human TSP-2 (SEQ ID NO: 2). The type 1 repeats of TSP-2 are, as illustrated here, 1) amino acids 381-436; 2) amino acids 437-493; and 3) amino acids 494-550.

Figure 3 is a representation of the amino acid sequence of human COMP (SEQ ID NO: 3). The type 2 repeats of COMP are, as illustrated here, 1) amino acids 89-128; 2) amino acids 129-181; 3) amino acids 182-226; and 4) amino acids 227-268

Figures 4A and 4B together are a representation of the DNA sequence (SEQ ID NO: 4) of gene encoding a human COMP/TSP-1 chimeric protein and the amino acid sequence (SEQ ID NO: 5) of a human COMP/TSP-1 chimeric protein encoded by the DNA sequence above it.

Figure 5A and 5B together are a representation of the DNA sequence (SEQ ID NO: 6) of a gene encoding a human COMP/TSP-2 chimeric protein and the amino acid sequence (SEQ ID NO: 7) of a human COMP/TSP-2 chimeric protein encoded by the DNA sequence above it.

Figure 6 is a schematic representation of a few of the chimeric protein embodiments of the invention.

Figure 7 is a graph showing tumor volume (mm³) at 7, 14 and 21 days in the experiment described in Example 3, in which mice were injected with an unaltered

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(control) vector, pNeo (filled diamonds) or with an expression vector encoding COMP/TSP-1 chimeric protein (filled squares).

DETAILED DESCRIPTION OF THE INVENTION

Described herein is a protein that has the functional activity of the TSR but not other activities associated with TSP-1 or TSP-2, and is assembled into a multimeric structure. One embodiment of the invention is a chimeric protein that comprises the TSRs from TSP-1 or TSP-2 and the multimer assembly region of human cartilage oligomeric matrix protein (COMP), using a portion of the aminoterminal end. Other portions of TSP-1 or TSP-2 can be incorporated into the chimeric protein, such as the procollagen homology region of TSP-1 and/or TSP-2. The last two TSRs of TSP-1 are preferably used because the first TSR has the ability to activate transforming growth factor β (TGF- β), which stimulates tumor growth. The COMP assembly domain spontaneously forms a 5-stranded α -helical domain, allowing for the use of the COMP domain as a tool for pentamerization.

Thus, the COMP/TSP-1 construct contains the region for multimerization, the first type 2 repeat of human COMP (construct encodes amino acids 1-128) and the second and third TSRs of human TSP-1 (construct encodes amino acids 417-530). See the Table for active sequences of TSP-1 (taken from chapter 2, "The Primary Structure of the Thrombospondins" In The Thrombospondin Gene Family 20 (J.C. Adams et al., eds.) Springer-Verlag, Heidelberg (1995)). The assembled protein is a pentamer containing 10 copies of the TSR. Thus, COMP/TSP-1 and COMP/TSP-2 are expected to be more active than TSP-1 and TSP-2. COMP/TSP-1 and COMP/TSP-2 are expected to be correctly folded and multimeric so that they better mimic the natural proteins than peptides that are based on the TSR sequence.

The first type 2 repeat of COMP includes amino acid residues 73-130, based on the genomic sequence. The amount of COMP sequence at the 3' end can be increased or decreased to maximize activity. For example, two or more type 2 repeats of COMP can be included if moving the type 1 repeats of TSP-1 or TSP-2 farther out on the arms of the expressed protein increases its activity. Alternatively, "spacer" sequence not naturally occurring in COMP or in TSP-1 or TSP-2 can be

added. The COMP/TSP-2 construct contains the same region of COMP and the three TSRs of human TSP-2 (construct encodes amino acids 381-550). When it is assembled to a pentamer this chimeric protein will contain 15 TSRs. Because these proteins are derived from portions of human proteins, they should not be immunogenic in humans.

Table: Active Regions of Interest Within Thrombospondin-1

Domain	Sequence	Function
Procollagen	NGVQYRN (SEQ ID NO: 8)	Anti-angiogenesis
homology		
Type 1 repeats	CSVTCG (SEQ ID NO: 9)	Cell binding
	WSXWSXW (SEQ ID NO: 10)	Heparin binding
	GGWSHW (SEQ ID NO: 11)	TGF-β and Fibronectin
		binding
	RFK	TGF-β activation
	SPWDICSVTCGGGVQKRSR	Anti-angiogenesis
	(SEQ ID NO: 12)	
Type 2 repeats	DVDEC(X) ₆ C(X) ₈ CENTDPGYNCLPC	Calcium binding
	(SEQ ID NO: 13)	

In one aspect, the invention comprises polynucleotides or nucleic acid molecules that encode chimeric proteins having portions whose amino acid sequences are derived from human TSP-1. By the genomic structure, the type 1 repeats of TSP-1 are amino acid residues 359-414 (first), amino acid residues 415-473 (second), and 474-531 (third). In one case, the chimeric protein encoded by the polynucleotides of the invention comprises the second and third type 1 repeats of human TSP-1. Such a chimeric protein may also comprise the procollagen homology region and the first type 1 repeat of TSP-1. If amino acid sequences that activate TGF-β are included in the product protein, and are found to reduce antiangiogenic activity, the RFK sequence can be mutated (to QFK, for example) to a

sequence that does not activate TGF- β , by appropriate manipulations of the nucleic acid molecule or construct encoding the chimeric proteins. In another case, the chimeric proteins encoded by the polynucleotides of the invention are variants of the immediately aforementioned chimeric protein which have activity that is similar in quality and quantity (for example, plus or minus one order of magnitude in an assay) to the anti-angiogenic activity of the protein whose amino acid sequence is represented in Figures 4A and 4B. In another case, the chimeric proteins encoded by polynucleotides of the invention comprise the second and third type 1 repeats of human TSP-1, the multimerization domain of human COMP, and the first type 2 repeat of human COMP. In another case, the chimeric proteins encoded by the polynucleotides of the invention are variants of the immediately aforementioned chimeric protein which have activity that is similar in quality and quantity to the anti-angiogenic activity of the protein whose amino acid sequence is represented in Figures 4A and 4B.

In one aspect, the invention comprises polynucleotides or nucleic acid molecules that encode chimeric proteins having portions whose amino acid sequences are derived from human TSP-2. The genomic structure of the human TSP-2 gene, which would provide one way to define the boundaries of the repeats, has not been determined. In one case, the chimeric protein encoded by the polynucleotides of the invention comprises the three type 1 repeats of human TSP-2. In another case, the chimeric proteins encoded by the polynucleotides of the invention are variants of the immediately aforementioned chimeric proteins which have activity that is similar in quality and quantity to the anti-angiogenic activity of the protein whose amino acid sequence is represented in Figures 5A and 5B. In another case, the chimeric protein encoded by polynucleotides of the invention comprises the three type 1 repeats of human TSP-2, and the multimerization domain of human COMP. In another case, the chimeric proteins encoded by the polynucleotides of the invention are variants of the immediately aforementioned chimeric protein which have activity that is similar in quality and quantity to the anti-angiogenic activity of the protein whose amino acid sequence is represented in Figures 5A and 5B.

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The polynucleotides of the invention can be made by recombinant methods, can be made synthetically, can be replicated by enzymes in *in vitro* (e.g., PCR) or *in vivo* systems (e.g., by suitable host cells, when inserted into a vector appropriate for replication within the host cells), or can be made by a combination of methods. The polynucleotides of the invention can include DNA and its RNA counterpart.

As used herein, "nucleic acid," "nucleic acid molecule," "oligonucleotide" and "polynucleotide" include DNA and RNA and chemical derivatives thereof, including phosphorothicate derivatives and RNA and DNA molecules having a radioactive isotope or a chemical adduct such as a fluorophore, chromophore or biotin (which can be referred to as a "label"). The RNA counterpart of a DNA is a polymer of ribonucleotide units, wherein the nucleotide sequence can be depicted as having the base U (uracil) at sites within a molecule where DNA has the base T (thymidine).

Isolated nucleic acid molecules or polynucleotides can be purified from a natural source or can be made recombinantly. Polynucleotides referred to herein as "isolated" are polynucleotides purified to a state beyond that in which they exist in cells. They include polynucleotides obtained by methods described herein, similar methods or other suitable methods, and also include essentially pure polynucleotides produced by chemical synthesis or by combinations of biological and chemical methods, and recombinant polynucleotides that have been isolated. The term "isolated" as used herein for nucleic acid molecules, indicates that the molecule in question exists in a physical milieu distinct from that in which it occurs in nature. For example, an isolated polynucleotide may be substantially isolated with respect to the complex cellular milieu in which it naturally occurs, and may even be purified essentially to homogeneity, for example as determined by agarose or polyacrylamide gel electorphoresis or by A_{260}/A_{280} measurements, but may also have further cofactors or molecular stabilizers (for instance, buffers or salts) added.

The invention further comprises the polypeptides encoded by the isolated nucleic acid molecules of the invention. Thus, for example, the invention relates to fusion proteins, comprising a portion of TSP-1 which comprises the second and third type 1 repeats, linked to a second moiety not occurring in TSP-1 as found in

nature. In an analogous manner, the invention relates also to fusion proteins, comprising TSP-2 or a functional portion thereof such as one or more repeat regions as a first moiety, linked to second moiety not occurring in TSP-2 as found in nature. The second moiety can be an amino acid, peptide or polypeptide, and can have enzymatic or binding activity of its own. The first moiety can be in an N-terminal location, C-terminal location or internal to the fusion protein. In one embodiment, the fusion protein comprises the portion of human TSP-1 described immediately above, or human TSP-2 or a portion thereof as the first moiety, and a second moiety comprising a linker sequence and an affinity ligand.

Another aspect of the invention relates to a method of producing a chimeric protein of the invention, or a variant thereof, and to expression systems and host cells containing a vector appropriate for expression of a chimeric protein of the invention. Variants of the chimeric protein include those having amino acid sequences that differ from those sequences in Figures 4A and 4B, and Figures 5A and 5B, wherein those variants have several, such as 5 to 10, 1 to 5, or 3, 2 or 1 amino acids substituted, deleted, or added, in any combination, compared to the sequences in Figures 4A and 4B and Figures 5A and 5B. In one embodiment, variants have silent substitutions, additions and deletions that do not alter the properties and activities of the chimeric protein. Variants can also be modified polypeptides in which one or more amino acid residues are modified, and mutants comprising one or more modified residues.

Proteins and polypeptides described herein can be assessed for their angiogenic activity by using an assay such as those described in Tolsma, S.S. et al., J. Cell Biol. 122(2):497-511 (1993), one which measures the migration of bovine adrenal capillary endothelial cells in culture, and one which tests migration of cells into a sponge containing an agent to be tested for activity. A further test for angiogenesis, which can also be adapted also to test anti-angiogenesis activity, is described in Polverini, P.J. et al., Methods. Enzymol. 198:440-450 (1991).

Cells that express such a chimeric protein or a variant thereof can be made and maintained in culture, under conditions suitable for expression, to produce protein for isolation. These cells can be procaryotic or eucaryotic. Examples of

procaryotic cells that can be used for expression (as "host cells"; "cell" including herein cells of tissues, cell cultures, cell strains and cell lines) include Escherichia coli, Bacillus subtilis and other bacteria. Examples of eucaryotic cells that can be used for expression include yeasts such as Saccharomyces cerevisiae,

Schizosaccharomyces pombe, Pichia pastoris and other lower eucaryotic cells, and cells of higher eucaryotes such as those from insects and mammals. Suitable cells of mammalian origin include primary cells, and cell lines such as CHO, HeLa, 3T3, BHK, COS, 293, and Jurkat cells. Suitable cells of insect origin include primary cells, and cell lines such as SF9 and High five cells. (See, e.g., Ausubel, F.M. et al., 10 eds. Current Protocols in Molecular Biology, Greene Publishing Associates and John Wiley & Sons Inc., (containing Supplements up through 1998)).

In one embodiment, host cells that produce a recombinant chimeric protein, variant, or portions thereof can be made as follows. A gene encoding a chimeric protein described herein can be inserted into a nucleic acid vector, e.g., a DNA vector, such as a plasmid, virus or other suitable replicon (including vectors suitable for use in gene therapy, such as those derived from adenovirus or others; see, for example Xu, M. et al., Molecular Genetics and Metabolism 63:103-109, 1998) can be present in a single copy or multiple copies, or the gene can be integrated in a host cell chromosome. A suitable replicon or integrated gene can contain all or part of the coding sequence for the protein or variant, operably linked to one or more expression control regions whereby the coding sequence is under the control of transcription signals and linked to appropriate translation signals to permit translation. The vector can be introduced into cells by a method appropriate to the type of host cells (e.g., transformation, electroporation, infection). For expression from the gene, the host cells can be maintained under appropriate conditions (e.g., in the presence of inducer, normal growth conditions, etc.). Proteins or polypeptides thus produced can be recovered (e.g., from the cells, the periplasmic space, culture medium) using suitable techniques.

The invention also relates to isolated proteins or polypeptides encoded by nucleic acids of the present invention. Isolated proteins can be purified from a natural source or can be made recombinantly. Proteins or polypeptides referred to

herein as "isolated" are proteins or polypeptides purified to a state beyond that in which they exist in cells and include proteins or polypeptides obtained by methods described herein, similar methods or other suitable methods, and also include essentially pure proteins or polypeptides, proteins or polypeptides produced by chemical synthesis or by combinations of biological and chemical methods, and recombinant proteins or polypeptides which are isolated. Thus, the term "isolated" as used herein, indicates that the polypeptide in question exists in a physical milieu distinct from the cell in which its biosynthesis occurs. For example, an isolated COMP/TSP-1 or COMP/TSP-2 chimeric protein may be purified essentially to homogeneity, for example as determined by PAGE or column chromatography (for example, HPLC), but may also have further cofactors or molecular stabilizers added to the purified protein to enhance activity. In one embodiment, proteins or polypeptides are isolated to a state at least about 75% pure; more preferably at least about 85% pure, and still more preferably at least about 95% pure, as determined by Coomassie blue staining of proteins on SDS-polyacrylamide gels.

Chimeric or fusion proteins can be produced by a variety of methods. For example, a chimeric protein can be produced by the insertion of a TSP gene or portion thereof into a suitable expression vector, such as Bluescript SK +/-(Stratagene), pGEX-4T-2 (Pharmacia), pET-15b, pET-20b(+) or pET-24(+) (Novagen). The resulting construct can be introduced into a suitable host cell for 20 expression. Upon expression, chimeric protein can be purified from a cell lysate by means of a suitable affinity matrix (see e.g., Current Protocols in Molecular Biology (Ausubel, F.M. et al., eds., Vol. 2, pp. 16.4.1-16.7.8, containing supplements up through Supplement 44, 1998).

Polypeptides of the invention can be recovered and purified from cell cultures by well-known methods. The recombinant protein can be purified by ammonium sulfate precipitation, heparin-Sepharose affinity chromatography, gel filtration chromatography and/or sucrose gradient ultracentrifugation using standard techniques. Further methods that can be used for purification of the polypeptide 30 include ethanol precipitation, acid extraction, anion or cation exchange chromatography, phosphocellulose chromatography, hydrophobic interaction

chromatography, affinity chromatography, hydroxylapatite chromatography and high performance liquid chromatography. Known methods for refolding protein can be used to regenerate active conformation if the polypeptide is denatured during isolation or purification.

5 The method to construct genes encoding COMP/TSP-1 or COMP/TSP-2 hybrid proteins can be applied more broadly to produce polynucleotides, and vectors and host cells comprising such polynucleotides, wherein the polynucleotides encode COMP/endostatin, COMP/angiostatin, COMP/platelet factor 4, or COMP/prolactin, for example. In each case, a portion of a polynucleotide known to encode full-length human endostatin, angiostatin, platelet factor 4 (GenBank Accession No. M25897) or prolactin (GenBank Accession No. V00566), can be chosen for cloning into a COMP cDNA as illustrated herein for COMP/TSP-1 and COMP/TSP-2 DNA constructs. Thus, the invention also includes COMP/endostatin, COMP/angiostatin, COMP/platelet factor 4, and COMP/prolactin chimeric proteins encoded by such nucleic acid constructs. See Figure 6 for a schematic representation of the structure of COMP/endostatin.

In addition, a portion of the endostatin, angiostatin, platelet factor 4 or prolactin coding regions, wherein that portion encodes a polypeptide having antiangiogenic activity, can be added to or incorporated into a DNA construct encoding COMP/TSP-1, such that a TSP-1-derived polypeptide and a polypeptide derived from endostatin, angiostatin, platelet factor 4 or prolactin are produced fused together in tandem on the same "arm" of the "5-armed" COMP-multimerized pentamer. Different expression constructs can be introduced into the same host cells such that two or more chimeric protein "arms" of different types (e.g.,

Chimeric protein antiangiogenic agents can be used, for example, after surgery or radiation to prevent recurrence of metastases, in combination with conventional chemotherapy, immunotherapy, or various types of gene therapy not necessarily directed against angiogenesis.

multimerization domain.

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Construction of COMP/TSP-1P Expression Vectors

Expression vectors that can be used to produce COMP/TSP-1P, a chimeric protein that includes the procollagen homology region (see Figure 6), can be produced from two distinct cDNAs. The COMP portion is identical to that in the 5 Examples described herein. For TSP-1, a new forward primer (GAT GAC GTC ACT GAA GAG AAC AAA GAG) (SEQ ID NO: 14) and the same reverse primer as described in the Examples can be used to produce a PCR product that is approximately 750 base pairs in size and has an AatII restriction endonuclease site at the 5' end and an Xbal restriction endonuclease site at the 3' end. The product codes 10 for amino acids 284-530 and includes the procollagen homology region (exons 6 and 7) and type 1 repeats. If inclusion of the TGF-B activating sequence (RFK) that is in the first type 1 repeat is found to reduce the antitumor activity, this sequence will be mutated to an inactive sequence (QFK, for example) using an oligonucleotidedirected mutagenesis kit (Amersham). The COMP/TSP-1P expression vector can be 15 constructed by cutting the PCR product with AatII and XbaI and cloning it into the COMP cDNA cut with the same enzymes. The protein can be expressed using the methods that have been described for COMP/TSP-1 and COMP/TSP-2.

Construction of COMP/Endostatin Expression Vectors

The strategy for making multimers of the TSP-1 and TSP-2 can be used to

20 make multimers of other anti-angiogenic proteins. For example, if the active region
of endostatin is prepared by PCR and cloned into the COMP cDNA, a pentameric
structure of endostatin can be made when this construct is expressed (O'Reilly M.D.,
et al., Cell 88:277-285, (1997)). In addition, if the COMP/TSP-1 and the
COMP/endostatin genes are expressed concurrently within the same cells, mixed

25 pentamers of COMP/TSP-1 and COMP/endostatin subunits are made. The mixed
multimer allows simultaneous treatment with the two reagents by delivery of a
single therapeutic. An additive or synergistic effect of the two agents may
significantly increase the efficacy of this reagent as compared to that of each reagent
alone. For example, combination therapy with angiostatin and endostatin has

eradicated tumors in mice (Boehm, T. et al., Nature 390:404-407, 1997).

The cDNA for endostatin can be prepared by PCR of liver cDNA or from an isolated cDNA clone for collagen XVIII (GenBank accession no. L22548). The human endostatin cDNA can be produced by PCR with the forward primer GAT GAC GTC CAC AGC CAC CGC G (SEQ ID NO: 15) and the reverse primer GAT 5 TCT AGA CTA CTT GGA GGC AGT CAT G (SEQ ID NO: 16). The resulting PCR product is approximately 560 base pairs and encodes amino acids 1 to 184 of human endostatin (Sasaki, T., et al., EMBO J., 17:4249-4256, 1998). The COMP/endostatin expression vector can be constructed by cutting the PCR product with AatII and XbaI, and cloning it into cDNA cut with the same enzymes. The 10 protein can be expressed using the methods that have been described herein for COMP/TSP-1 and COMP/TSP-2. Angiostatin, as it was isolated from mice bearing Lewis lung carcinoma, includes the first four kringle domains of plasminogen (amino acids 98-440) (O'Reilly, M.S., et al., Cell 79:315-328, 1994). It should be noted that smaller constructs that contain fewer kringle domains should also be active based on published data (Griscelli, F., et al., Proc. Natl. Acad. Sci. USA 95:6367-6372, 1998). A 16,000 dalton fragment of prolactin and platelet factor 4 have also been reported to inhibit angiogenesis (Clapp, C. et al., Endocrinology 133:1292-1299, 1993; Gapta, S.K., et al., Proc. Natl. Acad. Sci. USA 92:7799-7803, 1995).

Also included in the inventions are compositions containing, as a biological ingredient, an anti-angiogenic chimeric protein, or a variant thereof to inhibit angiogenesis in mammalian tissues, and use of such compositions in the treatment of diseases and conditions characterized by, or associated with, angiogenic activity. Such methods can involve administration by oral, topical, injection, implantation, 25 sustained release, or other delivery methods that bring one or more anti-angiogenic chimeric proteins in contact with cells whose growth is to be inhibited.

The present invention includes a method of treating an angiogenesismediated disease with a therapeutically effective amount of one or more antiangiogenic chimeric proteins. Angiogenesis-mediated diseases can include, but are 30 not limited to, cancers, solid tumors, tumor metastasis, benign tumors (e.g., hemangiomas, acoustic neuromas, neurofibromas, trachomas, and pyogenic

granulomas), rheumatoid arthritis, psoriasis, ocular angiogenic diseases (e.g., diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis), Osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularization, telangiectasia, hemophiliac joints, angiofibroma, and wound granulation.

"Cancer" means neoplastic growth, hyperplastic or proliferative growth or a pathological state of abnormal cellular development and includes solid tumors, nonsolid tumors, and any abnormal cellular proliferation, such as that seen in leukemia. As used herein, "cancer" also means angiogenesis-dependent cancers and tumors, 10 i.e., tumors that require for their growth (expansion in volume and/or mass) an increase in the number and density of the blood vessels supplying them with blood. "Regression" refers to the reduction of tumor mass and size. As used herein, the term "therapeutically effective amount" means the total amount of each active component of the composition or method that is sufficient to show a meaningful benefit to a treated human or other mammal, i.e., treatment, healing, prevention or amelioration of the relevant medical condition, or an increase in rate of treatment, healing, prevention or amelioration of such conditions. More specifically, for example, a therapeutically effective amount of an anti-angiogenic chimeric protein can cause a measurable reduction in the size or numbers of tumors, or in their rate of growth or multiplication, compared to untreated tumors. Other methods of assessing a "therapeutically effective amount," can include the result that blood vessel formation is measurably reduced in treated tissues compared to untreated tissues.

One or more anti-angiogenic chimeric proteins may be used in combination with other compositions and procedures for the treatment of diseases. For example, a tumor may be treated conventionally with surgery, radiation, chemotherapy, or immunotherapy, combined with anti-angiogenic chimeric proteins, and then anti-angiogenic chimeric proteins may be subsequently administered to the patient to extend the dormancy of micrometastases and to stabilize and inhibit the growth of any residual primary tumor.

The compositions may further contain other agents which either enhance the activity of the protein or compliment its activity or use in treatment, such as

chemotherapeutic or radioactive agents. Such additional factors and/or agents may be included in the composition to produce a synergistic effect with protein of the invention, or to minimize side effects. Additionally, administration of the composition of the present invention may be administered concurrently with other therapies, e.g., administered in conjunction with a chemotherapy, immunotherapy or radiation therapy regimen.

The angiogenesis-modulating composition of the present invention may be a solid, liquid or aerosol and may be administered by any known route of administration. Examples of solid compositions include pills, creams, and implantable dosage units. The pills may be administered orally, the therapeutic creams may be administered topically. The implantable dosage unit may be administered locally, for example at a tumor site, or may be implanted for systemic release of the angiogenesis-modulating composition, for example subcutaneously. Examples of liquid composition include formulations adapted for injection subcutaneously, intravenously, intraarterially, and formulations for topical and intraocular administration. Examples of aerosol formulation include inhaler formulation for administration to the lungs.

The anti-angiogenic chimeric proteins can be provided as isolated and substantially purified proteins in pharmaceutically acceptable formulations

(including aqueous or nonaqueous carriers or solvents) using formulation methods known to those of ordinary skill in the art. These formulations can be administered by standard routes. In general, the combinations may be administered by the topical, transdermal, intraperitoneal, intracranial, intracerebroventricular, intracerebral, intravaginal, intrauterine, oral, rectal or parenteral (e.g., intravenous, intraspinal, subcutaneous or intramuscular) route. In addition, the anti-angiogenic chimeric proteins may be incorporated into biodegradable polymers allowing for sustained release of the compound, the polymers being implanted in the vicinity of where drug delivery is desired, for example, at the site of a tumor, or implanted so that the anti-angiogenic chimeric proteins is slowly released systemically. Osmotic minipumps may also be used to provide controlled delivery of high concentrations of anti-angiogenic chimeric proteins through cannulae to the site of interest, such as directly

into a growth or into the vascular supply to that growth. The biodegradable polymers and their use are described, for example, in detail in Brem et al. (1991) (J. Neurosurg. 74:441-446), which is hereby incorporated by reference in its entirety.

As used herein, the terms "pharmaceutically acceptable," as it refers to compositions, carriers, diluents and reagents, represents that the materials are capable of administration to or upon a mammal with a minimum of undesirable physiological effects such as nausea, dizziness, gastric upset and the like. The preparation of a pharmacological composition that contains active ingredients dissolved or dispersed therein is well understood in the art and need not be limited based on formulation. Typically, such compositions are prepared as injectables either as liquid solutions or suspensions, however, solid forms suitable for solution, or suspensions, in liquid prior to use can also be prepared. The preparation can also be emulsified, for example, in liposomes.

The dosage of the anti-angiogenic chimeric proteins of the present invention will depend on the disease state or condition being treated and other clinical factors such as weight and condition of the human or animal and the route of administration of the compound. It is to be understood that the present invention has application for both human and veterinary use. The methods of the present invention contemplate single as well as multiple administrations, given either simultaneously or over an extended period of time.

The present invention also encompasses gene therapy whereby a polynucleotide encoding one or more anti-angiogenic chimeric proteins or one or more variants thereof, is introduced and regulated in a patient. Various methods of transferring or delivering DNA to cells for expression of the gene product protein, otherwise referred to as gene therapy, are disclosed in *Gene Transfer into Mammalian Somatic Cells in Vivo*, N. Yang (1992) *Crit. Rev. Biotechnol.* 12(4):335-356, which is hereby incorporated by reference. Gene therapy encompasses incorporation of DNA sequences into somatic cells or germ line cells for use in either ex vivo or in vivo therapy. Gene therapy can function to replace genes, augment normal or abnormal gene function, and to combat infectious diseases and other pathologies.

Strategies for treating these medical problems with gene therapy include therapeutic strategies such as identifying the defective gene and then adding a functional gene to either replace the function of the defective gene or to augment a slightly functional gene; or prophylactic strategies, such as adding a gene for the 5 product protein that will treat the condition or that will make the tissue or organ more susceptible to a treatment regimen. For example, a gene encoding an antiangiogenic chimeric protein may be inserted into tumor cells of a patient and thus inhibit angiogenesis.

Gene transfer methods for gene therapy fall into three broad categories: 10 physical (e.g., electroporation, direct gene transfer and particle bombardment), chemical (e.g., lipid-based carriers, or other non-viral vectors) and biological (e.g., virus-derived vector and receptor uptake). For example, non-viral vectors may be used which include liposomes coated with DNA. Such liposome/DNA complexes may be directly injected intravenously into the patient. It is believed that the 15 liposome/DNA complexes are concentrated in the liver where they deliver the DNA to macrophages and Kupffer cells. These cells are long lived and thus provide long term expression of the delivered DNA. Additionally, vectors or the "naked" DNA of the gene may be directly injected into the desired organ, tissue or tumor for targeted delivery of the therapeutic DNA.

In vivo gene transfer involves introducing the DNA into the cells of the patient when the cells are within the patient. Methods include using virally mediated gene transfer using a noninfectious virus to deliver the gene in the patient or injecting naked DNA into a site in the patient and the DNA is taken up by a percentage of cells in which the gene product protein is expressed. Additionally, the other methods described herein, such as use of a "gene gun," may be used for in vitro insertion of anti-angiogenic chimeric proteins DNA or anti-angiogenic chimeric proteins regulatory sequences.

Chemical methods of gene therapy may involve a lipid based compound, not necessarily a liposome, to transfer the DNA across the cell membrane. Lipofectins 30 or cytofectins, lipid-based positive ions that bind to negatively charged DNA, make a complex that can cross the cell membrane and provide the DNA into the interior of WO 00/44908 PCT/US00/02482

-18-

the cell. Another chemical method uses receptor-based endocytosis, which involves binding a specific ligand to a cell surface receptor and enveloping and transporting it across the cell membrane. The ligand binds to the DNA and the whole complex is transported into the cell. The ligand gene complex is injected into the blood stream and then target cells that have the receptor will specifically bind the ligand and transport the ligand-DNA complex into the cell.

Many gene therapy methodologies employ viral vectors to insert genes into cells. For example, altered retrovirus vectors have been used in *ex vivo* methods to introduce genes into peripheral and tumor-infiltrating lymphocytes, hepatocytes, epidermal cells, myocytes, or other somatic cells. These altered cells are then introduced into the patient to provide the gene product from the inserted DNA.

Viral vectors have also been used to insert genes into cells using in vivo protocols. To direct the tissue-specific expression of foreign genes, cis-acting regulatory elements or promoters that are known to be tissue-specific can be used.

Alternatively, this can be achieved using in situ delivery of DNA or viral vectors to specific anatomical sites in vivo. For example, gene transfer to blood vessels in vivo was achieved by implanting in vitro transduced endothelial cells in chosen sites on arterial walls. The virus infected surrounding cells which also expressed the gene product. A viral vector can be delivered directly to the in vivo site, by a catheter for example, thus allowing only certain areas to be infected by the virus, and providing long-term, site specific gene expression. In vivo gene transfer using retrovirus vectors has also been demonstrated in mammary tissue and hepatic tissue by injection of the altered virus into blood vessels leading to the organs.

Viral vectors that have been used for gene therapy protocols include but are not limited to, retroviruses, other RNA viruses such as poliovirus or Sindbis virus, adenovirus, adeno-associated virus, herpes viruses, SV40, vaccinia and other DNA viruses. Replication-defective murine retroviral vectors have been widely utilized gene transfer vectors.

Carrier mediated gene transfer *in vivo* can be used to transfect foreign DNA into cells. The carrier-DNA complex can be conveniently introduced into body fluids or the bloodstream and then site-specifically directed to the target organ or

tissue in the body. Both liposomes and polycations, such as polylysine, lipofectins or cytofectins, can be used. Liposomes can be developed which are cell specific or organ specific and thus the foreign DNA carried by the liposome will be taken up by target cells. Injection of immunoliposomes that are targeted to a specific receptor on certain cells can be used as a convenient method of inserting the DNA into the cells bearing the receptor. Another carrier system that has been used is the asialoglycoprotein/polylysine conjugate system for carrying DNA to hepatocytes for in vivo gene transfer.

The gene therapy protocol for transfecting anti-angiogenic chimeric proteins
into a patient may either be through integration of a gene encoding an antiangiogenic chimeric protein into the genome of the cells, into minichromosomes or
as a separate replicating or non-replicating DNA construct in the cytoplasm or
nucleoplasm of the cell. Anti-angiogenic chimeric proteins expression may continue
for a long-period of time or may be reinjected periodically to maintain a desired
level of the anti-angiogenic chimeric proteins protein in the cell, the tissue or organ
or a determined blood level.

EXAMPLES

Example 1: Construction of COMP/TSP-1 and COMP/TSP-2

The chimeric expression vectors have been produced from three distinct cDNAs. The first is a clone for human cartilage oligomeric matrix protein (COMP) and was isolated from a λgtll chondrocyte cDNA library (Doege, K.J, et al., J. Biol. Chem. 266:894-902 (1991)). This is an almost full-length clone for the COMP mRNA that only lacks a small region of the 5'-untranslated region. This clone (hCOMP-95) was used previously to determine the sequence of human COMP (GenBank Accession No. L32137; Genomics, 24:435-439 (1994)).

The second cDNA was produced using the polymerase chain reaction (PCR) with the human thrombospondin-1 (TSP-1) gene as the template. The TSP-1 clones were isolated from a human endothelial cell library (*J. Cell Biol. 103*:1635-1648

(1986)). The forward primer (GAT GAC GTC GAT GGT GGC TGG AGC CAC)
(SEQ ID NO: 17) and the reverse primer (GAT CTA GAT TGG ACA GTC CTG CTT G) (SEQ ID NO: 18) produce a PCR product that is approximately 354 basepairs in size and has an Aat II restriction endonuclease site at the 5' end and an
5 Xba I restriction endonuclease site at the 3' end. The PCR product encodes amino acids 417 to 530 and includes the second and third type 1 repeats of TSP-1 (see Figure 1 for the numbering of amino acids in TSP-1). The coding sequence for the first type 1 repeat was not included in the PCR product, by design, because it contains an RFK sequence that has been shown to activate TGF-β. This activity is not required to inhibit angiogenesis and it may produce unwanted secondary effects on numerous cell types. Vectors that include the first type 1 repeat can be constructed, using the same approach, if this region is found to enhance the antiangiogenic activity or other activities.

15 (catalog no. 936208 from Stratagene, LaJolla, CA) as the template. The forward primer (GAT GAC GTC GAG GAG GGC TGG TCT CCG) (SEQ ID NO: 19) and the reverse primer (GAT CTA GAC ACG GGG CAG CTC CTC TTG) (SEQ ID NO: 20) produced a PCR product that is approximately 520 base pairs in size and has an Aat II restriction endonuclease site at the 5' end and an Xba I restriction endonuclease site at the 5' end and an Xba I restriction of TSP-2 and, includes all three type 1 repeats of TSP-2 (see Figure 2 for numbering of amino acids in TSP-2). The sequence of the PCR primers was based on the human TSP-2 sequence in the GenBank database (Accession No. L12350). The sequences of the PCR products were determined to establish that mutations that affect the amino acid sequence had not been introduced during the PCR.

The COMP/TSP-1 and COMP-TSP-2 expression vectors were constructed by cutting the PCR products with Aat II and Xba I and subcloning them into the COMP cDNA vector [derived from Bluescript (Stratagene, La Jolla, CA)] cut with the same enzymes. The portion of COMP that was retained includes the signal sequence, the regions required for pentamerization and the first type 2 repeat (amino acids 1 to 128 on the enclosed sequence; Figure 3). Since there was an internal Aat

II site in the TSP-2 PCR product, it had to be cloned into the vector in two steps. A 430 basepair Aat II/Xba I fragment of the TSP-2 PCR product was subcloned into the vector containing the portion of COMP as a first step. The resulting subclone was cut with Aat II, and a 90 base pair Aat II fragment of the PCR product was 5 ligated into the expression vector. The final forms of the cDNAs were confirmed to have the predicted structure by nucleotide sequencing. They were then cut with Eco R1 and Xba I and ligated into the pcDNA 3.1 (Invitrogen; Carlsbad, CA) vector cut with the same enzymes. The DNA sequences of COMP/TSP-1 and COMP/TSP-2 are shown in Figures 4A and 4B and Figures 5A and 5B, respectively. The 10 predicted molecular weights of the subunits of COMP/TSP-1 and COMP/TSP-2 should be approximately 24,200 and 30,000, respectively. The fully assembled COMP/TSP-1 and COMP/TSP-2 proteins should be 121,000 Da and 150,000 Da, respectively. The amino acid sequences of these proteins are shown in Figures 4A and 4B and Figures 5A and 5B, respectively.

15 Example 2: Production of Isolated COMP/TSP-1 and COMP/TSP-2

To express these chimeric proteins, the expression vectors can be transfected into human kidney 293 cells using the Lipofectin protocol (Gibco Laboratories). The cells can be selected with Zeocin and individual clones can be grown. The secretion of COMP/TSP-1 and COMP/TSP-2 can be monitored with western 20 blotting using polyclonal antibodies to the region of COMP that is present in both expressed proteins. These antibodies have been produced by immunizing rabbits with a synthetically produced peptide, having an amino acid sequence derived from the N-terminal end of COMP, linked to a carrier protein. The amino acid sequence of the peptide is: SDLGPQMLRELQETN (SEQ ID NO: 21). A clone that 25 expresses high levels of the protein can be grown in large volume flasks and in serum free media.

Example 3: Inhibition of Tumor Growth by COMP/TSP-1

A cDNA of thrombospondin-1 (TSP-1) containing the second and third type-1 repeats and the COMP assembly sequence (COMP/TSP-1) was produced by PCR

using constructs derived as above as template, and was cloned into the expression vector pNeo (Invitrogen, Carisbad, CA). Both the resulting COMP/TSP-1 construct and the unaltered vector alone were transfected into the human squamous carcinoma cell line A431 (Streit, M., et al., American Journal of Pathology 155:441-452, 1999), and positive clones were selected using Geneticin at a concentration of 800 µg/ml. The growth curves of positive clones were determined over an 8 day period. Clones of pNeo- and COMP/TSP-1 construct-transfected cells that had similar growth curves were selected to test the effect of the chimeric protein on tumor growth in nude mice. A total of five mice pre group were injected intradermally at the shoulders with 5 X 10⁶ cells per site, two sites per mouse. Every week the tumors were measured with calipers. At three weeks, the mice were sacrificed and the tumors were removed for further studies. As can be seen from Figure 7, expression of COMP/TSP-1 caused inhibition of the growth of the tumors in this model.

All references (e.g., journal articles, books, published patent applications and patents, etc.) cited herein are hereby incorporated by reference.

While this invention has been particularly shown and described with references to preferred embodiments thereof, it will be understood by those skilled in the art that various changes in form and details may be made therein without departing from the spirit and scope of the invention as defined by the appended claims.

15

CLAIMS

What is claimed is:

- A nucleic acid molecule encoding a chimeric protein comprising the second and third type 1 repeats of human TSP-1, but not the TGF-β activation region of human TSP-1.
- A nucleic acid molecule encoding a chimeric protein comprising the
 multimerization domain of human COMP, the first type 2 repeat of human
 COMP, and the second and third type 1 repeats of human TSP-1.
- A nucleic acid molecule encoding a chimeric protein comprising the
 multimerization domain of human COMP, the first type 2 repeat of human
 COMP, and the second and third type 1 repeats of human TSP-1, but not the
 TGF-β activation region of human TSP-1.
 - 4. A nucleic acid molecule encoding a chimeric protein comprising the multimerization domain of human COMP, the procollagen homology region of TSP-1, and the first, second, and third type 1 repeats of human TSP-1.
 - 5. A nucleic acid molecule encoding a chimeric protein comprising the multimerization domain of human COMP, the procollagen homology region of TSP-1, and the first, second, and third type 1 repeats of human TSP-1, but not the TGF- β activation region of human TSP-1.
- 20 6. A nucleic acid molecule encoding a chimeric protein comprising the multimerization domain of human COMP and a portion of human endostatin, wherein the chimeric protein has anti-angiogenic activity.

- -24-
- A nucleic acid molecule encoding a chimeric protein comprising the 7. multimerization domain of human COMP and a portion of human angiostatin, wherein the chimeric protein has anti-angiogenic activity.
- A nucleic acid molecule encoding a chimeric protein comprising the 8. multimerization domain of human COMP and a portion of human prolactin, 5 wherein the chimeric protein has anti-angiogenic activity.
 - A nucleic acid molecule encoding a chimeric protein comprising the 9. multimerization domain of human COMP and a portion of human platelet factor 4, wherein the chimeric protein has anti-angiogenic activity.
- A nucleic acid molecule encoding a chimeric protein comprising the 10. 10 multimerization domain of human COMP, the procollagen homology region, and the first, second, and third type 1 repeats of human TSP-1.
 - A nucleic acid molecule encoding a protein having the amino acid sequence 11. SEQ ID NO: 5.
- A vector comprising nucleic acid encoding a chimeric protein comprising the 15 12. second and third type 1 repeats of human TSP-1 but not the TGF- β activation region of human TSP-1.
 - A host cell comprising the vector of Claim 12. 13.
- A vector comprising nucleic acid encoding a chimeric protein comprising the 14. multimerization domain of human COMP, the first type 2 repeat of human 20 COMP, and the second and third type 1 repeats of human TSP-1.
 - A host cell comprising the vector of Claim 14. 15.

- 16. A method for producing a chimeric protein which comprises the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the second and third type 1 repeats of human TSP-1, said method comprising maintaining the host cell of Claim 15 under conditions suitable for expression of said nucleic acid, whereby said protein is produced.
- 17. The method of Claim 16 further comprising isolating the chimeric protein.
- A vector comprising nucleic acid encoding a chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human
 COMP, and the second and third type 1 repeats of human TSP-1, but not the TGF-β activation region of human TSP-1.
 - 19. A host cell comprising the vector of Claim 18.
- A method for producing a chimeric protein which comprises the multimerization domain of human COMP, the first type 2 repeat of human
 COMP, and the second and third type 1 repeats of human TSP-1, but not the TGF-β activation region of human TSP-1, said method comprising maintaining the host cell of Claim 19 under conditions suitable for expression of said nucleic acid, whereby said protein is produced.
 - 21. The method of Claim 20 further comprising isolating the chimeric protein.
- 20 22. A vector comprising nucleic acid encoding a chimeric protein comprising the multimerization domain of human COMP, the procollagen homology region, and the first, second, and third type 1 repeats of human TSP-1.
 - 23. A vector comprising nucleic acid encoding a protein having the amino acid sequence SEQ ID NO: 5.

- 24. A host cell comprising the vector of Claim 23.
- 25. A chimeric protein comprising the second and third type 1 repeat of human TSP-1, but not the TGF- β activation region of human TSP-1.
- A chimeric protein comprising the multimerization domain of human
 COMP, the first type 2 repeat of human COMP, and the second and third type 1 repeats of human TSP-1.
 - 27. A chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the second and third type 1 repeats of human TSP-1, but not the TGF-β activation region of human TSP-1.
 - 28. A chimeric protein comprising the multimerization domain of human COMP, the procollagen homology region of TSP-1, and the first, second, and third type 1 repeats of human TSP-1.
- A chimeric protein comprising the multimerization domain of human COMP
 and a portion of human endostatin, wherein the chimeric protein has antiangiogenic activity.
 - 30. A chimeric protein comprising the multimerization domain of human COMP and a portion of human angiostatin, wherein the chimeric protein has antiangiogenic activity.
- 20 31. A chimeric protein comprising the multimerization domain of human COMP and a portion of human prolactin, wherein the chimeric protein has antiangiogenic activity.

- 32. A chimeric protein comprising the multimerization domain of human COMP and a portion of human platelet factor 4, wherein the chimeric protein has anti-angiogenic activity.
- 33. A protein having the amino acid sequence SEQ ID NO: 5.
- 5 34. An isolated nucleic acid molecule encoding a chimeric protein comprising the three type 1 repeats of human TSP-2.
 - 35. A vector comprising nucleic acid encoding a chimeric protein comprising the three type 1 repeats of human TSP-2.
 - 36. A host cell comprising the vector of Claim 35.
- 10 37. A method for producing a chimeric protein which comprises the three type 1 repeats of human TSP-2, said method comprising maintaining the host cell of Claim 36 under conditions suitable for expression of said nucleic acid, whereby said protein is produced.
 - 38. The method of Claim 37 further comprising isolating the chimeric protein.
- 15 39. A nucleic acid molecule encoding a chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the three type 1 repeats of human TSP-2.
- 40. A vector comprising isolated nucleic acid encoding a chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the three type 1 repeats of human TSP-2.
 - 41. A host cell comprising the vector of Claim 40.

- 42. A method for producing a chimeric protein which comprises the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the three type 1 repeats of human TSP-2, said method comprising maintaining the host cell of Claim 41 under conditions suitable for expression of said nucleic acid, whereby said protein is produced.
- 43. The method of Claim 42 further comprising isolating the chimeric protein.
- 44. A nucleic acid molecule encoding a protein having the amino acid sequence SEQ ID NO: 7.
- 45. A vector comprising nucleic acid encoding a protein having the amino acid sequence SEQ ID NO: 7.
 - 46. A host cell comprising the vector of Claim 45.
 - 47. A chimeric protein comprising the three type 1 repeats of human TSP-2.
 - 48. A chimeric protein comprising the procollagen homology region of TSP-2 and the three type 1 repeats of human TSP-2.
- 15 49. A chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the three type 1 repeats of human TSP-2.
 - 50. A protein having the amino acid sequence SEQ ID NO: 7.
- A method for inhibiting angiogenesis in a human or other mammal, the method comprising administering to the human or other mammal a therapeutically effective amount of an anti-angiogenic chimeric protein.

- 52. The method of Claim 51 wherein the anti-angiogenic chimeric protein is selected from the group consisting of:
 - a) a chimeric protein comprising the second and third type 1 repeats of human TSP-1;
- b) a chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the second and third type 1 repeats of human TSP-1;
 - c) a chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the second and third type 1 repeats of human TSP-1, but not the TGF-β activation region of human TSP-1;
 - d) a chimeric protein comprising the multimerization domain of human COMP, the procollagen region, and the first, second, and third type 1 repeats of human TSP-1; and
- a chimeric protein comprising the three type 1 repeats of human TSP-2; and (6) a chimeric protein comprising the multimerization domain of human COMP, the first type 2 repeat of human COMP, and the three type 1 repeats of human TSP-2.
- 53. The method of Claim 51 wherein the anti-angiogenic protein is administered locally at the site of one or more growths.

-	human t	thrombospondin-1
NH ₂	1 61 121 181	HRIPESGGDHSVFDIFELTGAARKGSGRRLVKGPDPSSPAFRIEDANLIPPVPDDKFQDL VDAVRTEKGFLLLASLRQMKKTRGTLLALERKDHSGQVFSVVSNGKAGTLDLSLTVQGKQ HVVSVEBALLATGQWKSITLFVQEDRAQLYIDCEKMENAELDVPIQSVFTRDLASIARLR JAKGGVNDHFQGVLQNVRFVFGTTPEDILRHKGCSSSTSVLLTLDHHVV(NGSSPAIRTHY
	2 1 1	IGHKTKDLQAICGISCDELSSM
۵	312	VLELRGLRTIVTTLQDSIRKVTEENKELANELRRPPLCYHHGVQYRNHE EMTVDSCTECHCQNSVTICKKVSCPINPCSÜÄATVPDGECCPRCMPSDSA
lype 1	361 417 474	DDGWSPWSEWTSCSTSCGHGIQQRGRSCDSLHNRCEGSSVQTRTCHIQECDKRFKQ DGGWSHWSPWSS <mark>CSVTCG</mark> DGVITRIRLCHSPSPQHGKPCEGEARETKACKKDACPI HGGWGPWSPWDICS <u>VTCG</u> GGVQKRSR ¹ LCHMPPPQFGGKDCVGDVTEHQICHKQDCPI
type 2	531 572 630	DGCLSNPCFAGVKCTSYPDGSWKCGACPPGYSGHG1QCTDV DECKEVPDACFNHNGEHRCENTDPGYHCLPCPPRFTGSQPFGQGVEHATAUKQVCKPR NPCTDGTHDCNKHAKCHYLGHYSDPMYRC-ECKFGYAGHG11CGE
	674	DTDLDGWPHEHLVCVAGNTYHCKK
(ype 3	698 734 757 793 816 854	DRCPULPHSGQEDYDKDGIGDACDDDDDHUKIPDDR DRCPFHYHPAQYDYDRDDVGBRC DRCPYHHPDQADTDHHGEGDACAADIDGBGILHER DRCQYYYHVDQRDTBHGGYGDACAADIDGBGILHER DRCPLEHRPUQLDSDSDRIGDTCHMQDIDEDGHQHHL DRCPLYPHANQADHDKDGKGDACDHDDDHUGIPDDK
СООН	926 986 1046 1106	, b dicpenvoisetoprrpomiplopkgtsquophwvvrhqgkelvqtvhcdpglavgydev s navdfsgtppinterdddyagfvfgyqsssr <u>pyvymmk</u> qvtqsywdt@pyrraqgysglsv s kvv(i)sttgpgehlrnalwhtghtpgqvrtlnhdprhigwkdftayrwrlshrpktgf <u>irv</u> s <u>vm</u> yegkkimadsgpiydktyaggrlglevfsqenvfpsdlkyecrdp
_		

FIG. 1

	human	thrombospondin-2
NH ₂	1 61 121 181	MVWRLVLLALWVWPSTQAGUQDKDTTFDLFSISNINRKTIGAKQFRGPDPGVPAYRFVRF DYIPPVNADDLSKITKIHRQKEGFFLTAQLKQDGKSRGTLLALEGPGLSQRQFEIVSNGP ADTLDLTYWIDGTRHVVSLEDVGLADSQWKÜNTVQVAGETYSLHVGCDLIGPVALDEPFY EHLQAEKSHMYVAKGSARESHFRGLLQHVHLVFENSVEDILSKKGCQQGGGAEINAISEN
	241	TET LIRLG PHVTTEYVG PSSERRPEVCERSCEELG HM
۵	332	VQELSGLHVLVNQLSENLKRVSNDHQFLWELIGGPPKTRÜMSACHQDGRFFALÜBE TWVVDSCTTCTCKKFKTICHQITCPPATCASPSFVEGECCPSCLHSVDG
type 1	381 437 494	EEGWSPWAEWTQCS <u>VTCG</u> SGTQQRGRSCDVTSNTCLGPSIQTRACSLSKCDTRIRQ DGGWSHWSPWSSCS <u>VTCG</u> VGŴITRIRLCHSPVPQMGGKHCKGSGRETKACQGAPCPI DGRWSPWSPWSACTVTCAGGIRERTRVCHSPEPQYGGKACVGDVQERQMCNKRSCPV
type 2	551 592 650	DGCLSHPCFPGAQCSSFPDGSHSCGFCPVGFLG
	694	В ВВЪВВИНИ БИСАТ(ЙАТУНСТК
lype 3		DHCPHLPHSGQEDFDKDGIGDACDDDDDNDGVTDEK DNCQLLFNPRQADYDKDEVGDRC DHCPYVHNPAQIDTDNNGEGDACSVDIDGDDVFNER DHCPYVHYDQRDTDGDGVGDHC DHCPLVHPPQTDVDNDLVGDQCDNNEDIDDDGHQHHQ DHCPLVHPPQTDVDNDLVGDQCDNNEDIDDDGHQHHQ
СООН	910 946 1006 11066	DRCRLVENPDQEDLDGDG <u>RGDI</u> CKDDFDNDN1PD1D DVCPENNAISETDFRNFQMVPLDPKGTTQIDPNWVIRHQGKELVQTANSDPGIAVGFDEF GSVDFSGTFYVNTDRDDDXAGFVFGYQSSSR <u>FYVVMMK</u> QVTQTYWEDQPTRAYGYSGVSL KVV@BSTTGTGEHLRNALHHTGNTPGQVRTLWHDPRHIGWKDYTAYRWHLTHRPKTGYIRV LVHEGKQVMADSGPIYDQTYAGGRLGLFVFSQEMVYFSDLKYECRDI

16, 2

518 DVCPENAEVTLTDFRAFQTVVLDPEGDAQIDPNWVVLNQGREIVQTMNSDPGLAVGYTAF HGVDFEGTFHVNTVTDDDYAGF1FGYQDSSS<u>FYVVMMK</u>QMEQTYWQANPFRAVAEPG1QL R F Y E G P E I. V A D S N V V L D T T M R G G R L G V F C F S Q E N I I W A N L R Y R C (i) D T I P E D Y E T H Q I. R Q A KAVKSSTGPGEQLRNALWHTGDTESQVRLLWKDPRNVGWKDKKSYRWFLQHRPQVGYIRV MVPDTACVLLLTLAALGASGQGQSPLGSDLGPQMLRELQETNAALQDVRDWLRQQVREIT -----DQASGCQRGAQ HECNAHP---CFPRVRCIGITSPGFR-CEACPPGYSGPTHQGVGLAFAKANKQVCTDI --NGILCGR 89 LHCAPGF --- CFPGVACIQTESGGR-CGPCPAGFTG--lype 2 182 NECETGOHN-CVPNSVCINTRGSFQ-CGPCQPGFVG---DNCPQKSNPDQADVDHDFVGDACDSDQDQDGDGHQDSR DNCPTVPNSAQEDSDHDGQGDACD--DDDDDNDGVPDSR DNCRLVPHPGQEDABRUGVGBVCQ---DDFDABKVVDKI DUCVTVPNSGQEDVDRUG1GDACD--PDAUGDGVPNEK DNCRSQKNDDQKDTDQDGRGD Δ CD...-DDTDGDRIRNQA RFCPDGSPSECHEHADCVI.ERDGSRSCV-CRVGWAG-61 FLKNTVMECDACGMQQSVRTGLPSVRPL DNCPLVRNPDQRNTDEDKWGDAC DNCPRVPNSDQKDSDGDG1GDAC DTDLDGFPDEKLRCPEPQCRK human COMP COOH 578 638 290 349 type 3 3 8 5 446 4 0 8 326 129

16.3

CAGCACCCAG CTCCCCGCCA CCGCC	ATG GTC	CCC GAC	ACC GCC TO	C GTT	CTT 52
	Met Val	Pro Asp	Thr Ala Cy 5	s Val	Leu
CTG CTC ACC CTG GCT GCC CTC Leu Leu Thr Leu Ala Ala Leu 10 15	GGC GCG Gly Ala	TCC GGA Ser Gly 20	CAG GGC CA Gln Gly Gl	G AGC n Ser	CCG 100 Pro 25
TTG GGC TCA GAC CTG GGC CCG Leu Gly Ser Asp Leu Gly Pro 30	CAG ATG	CTT CGG : Leu Arg 35	GAA CTG CA Glu Leu Gl	G GAA . n Glu 40	ACC 148 Thr
AAC GCG GCG CTG CAG GAC GTG Asn Ala Ala Leu Gln Asp Val 45	CGG GAC Arg Asp	Trp Leu	Ard GTU GT	G GTC n Val	AGG 196 Arg
GAG ATC ACG TTC CTG AAA AAC Glu Ile Thr Phe Leu Lys Asn 60	ACG GTG Thr Val	ATG GAG L Met Glu	TGT GAC GG Cys Asp Al	G TGC La Cys	GGG 244 Gly
ATG CAG CAG TCA GTA CGC ACC Met Gln Gln Ser Val Arg Thr 75 80	Gly Lev	A CCC AGC 1 Pro Ser	GTG CGG CG Val Arg Pi 85	CC CTG	CTC 292 Leu
CAC TGC GCG CCC GGC TTC TGC His Cys Ala Pro Gly Phe Cys 90 95	TTC CCC	GGC GTG Gly Val 100	GCC TGC A	rc CAG le Glm	ACG 340 Thr 105
GAG AGC GGC GGC CGC TGC GGC Glu Ser Gly Gly Arg Cys Gly 110	CCC TGG	C CCC GCG s Pro Ala 115	GGC TTC AGGly Phe Ti	og ggc nr gly 120	AAC 388 Asn

FIG. 4A

		Thr							CCG Pro	436
									ATC Ile	484
				AGC Ser 160					GAA Glu	532
				TAY						580
				TGG Trp						628
				AAA Lys					ACA Thr	676
				GAC Asp						724
			Asp	TGT Cys 240	 	 A				755

FIG. 4B

CAGC	ACC(CAG (CTCC	CCGC	CA C	ZGCC	ATG Met 1	GTC Val	CCC Pro	GAC Asp	ACC Thr 5	GCC Ala	TGC Cys	GTT Val	CTT Leu	52
CTG Leu 10	CTC Leu	ACC Thr	CTG Leu	GCT Ala	GCC Ala 15	CTC Leu	GGC Gly	GCG Ala	TCC Ser	GGA Gly 20	CAG Gln	GGC Gly	CAG Gln	AGC Ser	CCG Pro 25	100
TTG Leu	GGC Gly	TCA Ser	GAC Asp	CTG Leu 30	GGC Gly	CCG Pro	CAG Gln	ATG Met	CTT Leu 35	CGG Arg	GAA Glu	CTG Leu	CAG Gln	GAA Glu 40	ACC Thr	148
AAC Asn	GCG Ala	GCG Ala	CTG Leu 45	CAG Gln	GAC Asp	GTG Val	CGG Arg	GAC Asp 50	TGG Trp	CTG Leu	CGG Arg	CAG Gln	CAG Gln 55	GTC Val	AGG Arg	196
GAG :	ATC Ile	ACG Thr 60	TTC Phe	CTG Leu	AAA Lys	AAC Asn	ACG Thr 65	GTG Val	ATG Met	GAG Glu	TGT Cys	GAC Asp 70	GCG Ala	TGC Cys	GGG Gly	244
ATG Met	CAG Gln 75	CAG Gln	TCA Ser	GTA Val	CGC Arg	ACC Thr 80	GGC Gly	CTA Leu	CCC Pro	AGC Ser	GTG Val 85	CGG Arg	CCC Pro	CTG Leu	CTC Leu	292
CAC 'His 90	TGC Cys	GCG Ala	CCC Pro	GGC Gly	TTC Phe 95	TGC Cys	TTC Phe	CCC Pro	GGC Gly	GTG Val 100	GCC Ala	TGC Cys	ATC Ile	CAG Gln	ACG Thr 105	340
GAG . Glu	AGC Ser	GGC Gly	GGC Gly	CGC Arg 110	TGC Cys	GGC Gly	CCC Pro	TGC Cys	CCC Pro 115	GCG Ala	GGC Gly	TTC Phe	ACG Thr	GGC Gly 120	AAC Asn	388
GGC Gly	TCG Ser	CAC His	TGC Cys 125	ACC Thr	GAC Asp	GTC Val	GAG Glu	GAG Glu 130	GGC Gly	TGG Trp	TCT Ser	CCG Pro	TGG Trp 135	GCA Ala	GAG Glu	436
TGG Trp	ACC Thr	CAG Gln 140	TGC Cys	TCC Ser	GTG Val	ACG Thr	TGT Cys 145	GGC Gly	TCT Ser	GGG Gly	ACC Thr	CAG Gln 150	CAG Gln	AGA Arg	GGC	484

FIG. 5A

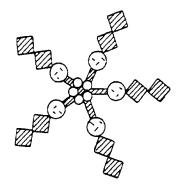
Arg	Ser 155	Cys	GAC Asp	Val	Thr	Ser 160	ASI	III	Суз	ne.	165					532
ACA Thr 170	CGG Arg	GCT Ala	TGC Cys	AGT Ser	CTG Leu 175	AGC Ser	AAG Lys	TGT Cys	GAC Asp	ACC Thr 180	CGC Arg	ATC Ile	CGG Arg	CAG Gln	GAC Asp 185	580
GGC Gly	GGC Gly	TGG Trp	AGC Ser	CAC His	TGG Trp	TCA Ser	CCT Pro	TGG Trp	TCT Ser 195	TCA Ser	TGC Cys	TCT Ser	GTG Val	ACC Thr 200	TGT Cys	623
GGA Gly	GTT Val	GGC Gly	AAT Asn 205	ATC Ile	ACA Thr	CGC Arg	ATC Ile	CGT Arg 210	CTC Leu	TGC Cys	AAC Asn	TCC	CCA Pro 215	GTG Val	CCC Pro	67 <i>6</i>
CAG Gln	ATG Met	GGG Gly 220	GGC Gly	AAG Lys	AAT Asn	TGC Cys	AAA Lys 225	GGG	AGT Ser	GGC Gly	CGG Arg	GAG Glu 230	ACC Thr	AAA Lys	GCC Ala	724
TGC Cys	CAG Gln 235	GGC Gly	GCC Ala	CCA Pro	TGC Cys	CCA Pro 240	ATC Ile	GAT Asp	GGC Gly	CGC Arg	TGG Trp 245		CCC Pro	TGG Trp	TCC Ser	772
CCG Pro 250	Trp	TCG Ser	GCC Ala	TGC Cys	ACT Thr 255	Val	ACC Thr	TGT Cys	GCC Ala	GGT Gly 260	Gry	ATC	CGG Arg	GAG Glu	CGC Arg 265	820
ACC Thr	CGG Arg	GTC Val	TGC Cys	AAC Asn 270	Ser	CCT Pro	GAG Glu	CCT Pro	CAG Gln 275	. TĀT	GGA Gly	GGG Gly	AAG Lys	GCC Ala 280	TGC Cys	8 € 8
GTG Val	GGG Gly	GAT Asp	GTG Val	Glr	GAG Glu	G CGI Arg	CAG	ATG Met 290	Cys	: AAC : Asr	AAC Lys	AGG Arg	AGC Ser 295	-1-	CCC Pro	916
	TCT Ser															925

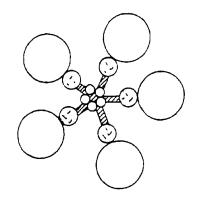
FIG. 5B

COMP/TSP-1

8/9

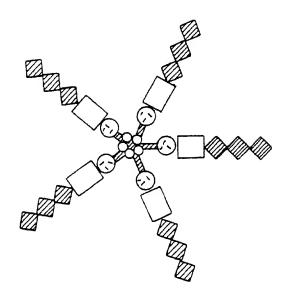
COMP/ENDOSTATIN

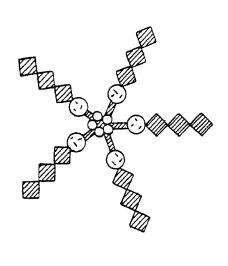




COMP/TSP-1P

COMP/TSP-2





pentamerization domain of human COMP

 \bigcirc

type 2 repeat of human COMP

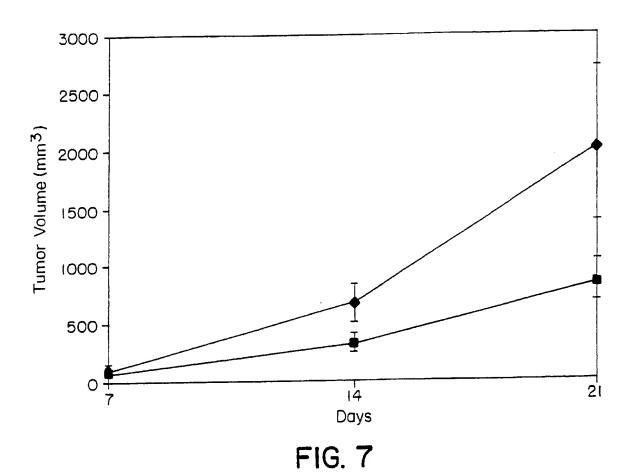
second and third type 1 repeats of TSP-1

all three type 1 repeats of TSP-1 or-2

procollagen homology region

endostatin

FIG. 6
SUBSTITUTE SHEET (RULE 26)



Intern 1al Application No PCT/US 00/02482

		PC1/US 00	7 02402
A. CLASS IPC 7	C12N15/52 C07K14/78 C07K1	9/00 A61K38/17	
According t	to International Patent Classification (IPC) or to both national clas	sification and IPC	
	SEARCHED		
Minimum di IPC 7	ocumentation searched (classification system followed by classification s	ication symbols)	
Documenta	ation searched other than minimum documentation to the extent the	nat such documents are included. In the fields se	earched
Electronic d	data base consulted during the international search (name of data	a base and, where practical, search terms used)
EPO-In Data	ternal, CHEM ABS Data, BIOSIS, MED	DLINE, SCISEARCH, BIOTECH	NOLOGY ABS, WPI
	ENTS CONSIDERED TO BE RELEVANT		
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4	1999, 274 (1) P430-7, XP002140107 UNITED STATES abstract; figure 2B	, UAIL I	2-5,10, 11,
		-/	14-24, 26-28
Y Furti	her documents are listed in the continuation of box C.	Patent family members are listed in	n annex.
Special co	tegories of cited documents :		
A* docume consid	ant defining the general state of the an which is not dered to be of particular relevance	Telater document published after the inter or priority date and not in conflict with to cited to understand the principle or the invention	the application but ory underlying the
filing d	document but published on or after the International late ant which may throw doubts on priority claim(s) or is cited to establish the publication date of another	"X" document of particular relevance; the cl cannot be considered novel or cannot involve an inventive step when the doc "Y" document of particular relevance; the cl	be considered to sument is taken alone
citation O° docume other r	n or other special reason (as specified) ent referring to an oral disclosure, use, exhibition or	cannot be considered to involve an inv document is combined with one or moi ments, such combination being obviou in the art.	entive step when the re other such docu-
later th	nan the priority date claimed	"&" document member of the same patent f	
	actual completion of the international search 1 September 2000	Date of mailing of the international sea	rch report
	mailing address of the ISA European Patent Office, P.B. 5818 Patentiaan 2	Authorized officer	
	NL – 2280 HV Rijswijk Tel. (+31-70) 340-2040, Tx. 31 851 epo nl. Fax: (+31-70) 340-3016	Gurdjian, D	

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C.(Continu	ation) DOCUMENTS CONSIDERED TO BE RELEVANT Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
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A	WO 98 39418 A (ARIAD GENE THERAPEUTICS INC; GILMAN MICHAEL Z (US)) 11 September 1998 (1998-09-11) claims 1-5	1-5, 10-28,33
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alegory °	,	Relevant to claim No.
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	O'ROURKE KM ET AL: "Thrombospondin 1 and thrombospondin 2 are expressed as both homo- and heterotrimers." JOURNAL OF BIOLOGICAL CHEMISTRY, DEC 15 1992, 267 (35) P24921-4, XP002147183 UNITED STATES the whole document	34-50

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Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet) This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons: Claims Nos because they relate to subject matter not required to be searched by this Authority, namely: Although claims 51-53 are directed to a method of treatment of the human/animal body, the search has been carried out and based on the alleged effects of the compound/composition. Claims Nos.: because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically: Claims Nos.: because they are dependent claims and are not drafted in accordance with the second and third sentences of Ruie 6.4(a). Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet) This International Searching Authority found multiple inventions in this international application, as follows: see additional sheet As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims. As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee. As only some of the required additional search fees were timely paid by the applicant, this international Search Report covers only those claims for which fees were paid, specifically claims Nos.: No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.: The additional search fees were accompanied by the applicant's protest. Remark on Protest No protest accompanied the payment of additional search fees.

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

This International Searching Authority found multiple (groups of) inventions in this international application, as follows:

1. Claims: 51-53 partially and 1-5,10-28 ,33

TSP-1 chimeric protein constructs, corresponding nucleic acids ,vectors ,host cells , method of producing chimeric proteins

2. Claims: 51-53 partially and 34-50

TSP-2 chimeric protein constructs, corresponding nucleic acids ,vectors ,host cells ,method of producing chimeric proteins

3. Claims: 51-53 partially and 6-9, 29-32

COMP chimeric protein constructs, not provided in the first and second invention of present application, corresponding nucleic acids, vectors, host cells, method of producing chimeric proteins

Information on patent family members

Intern and Application No PCT/US 00/02482

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